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GenCore version 4.5
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- protein search, using sw model OM protein

February 5, 2001, 12:02:32; search time 34.92 Seconds (without alignments) 7.778 Million cell updates/sec Run on:

US-09-687-267-6

1 XXXC 4 Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

195891 seqs, 67900655 residues Searched:

195891 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

PIR_66:* Database :

1: pirl: * 2: pir2: * 3: pir3: * 4: pir4: *

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

Ig H chain V-D-J r T-cell receptor be hypothetical L2 pr Ig heavy chain V r metallothionein is urine glycopeptide L-serine dehydrata leucinetRNA liga acylase - Kluyvera R-phycoerythrin ga variant surface gl variant surface gl variant surface gl rariant surface gl	T-cell receptor be
PH1602 PH0932 15695 15695 S9622 XGHUEU A25836 PC1002 S19288 A37521 C61512 D61512 PH1618 PH1618	PH0934
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ALIGNMENTS

SULT 1	R-phycoerythrin alpha-1 chain - red alga (Gastroclonium coulteri) (fragment) C:Species: Gastroclonium coulteri	C:Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993 C:Accession: A22565	R;Klotz, A.V.; Glazer, A.N. J. Biol. Chem. 260, 4856-4863, 1985	A; Title: Characterization of the bilin attachment sites in R-phycoerythrin. A; Reference number: A22565; MUID:85182601	A; Accession: A22565	A;Molecule type: protein A;Residues: 1-3 <klo></klo>
RESULT A22565	R-phyc	C; Date	R; Klot	A; Titl A; Refe	A; Acce	A;Mole A;Resi

ö Gaps ; 0 0; Indels Length 3; DB 2; . 0; Query Match 75.0%; Score 9; DB 2 Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches

4 C 4 - U ò g

metallothionein-A - rainbow trout (fragment)
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Accession: 151049
R;Olsson, P.E.; Kling, P.; Erkell, L.J.; Kille, P.
R;Olsson, P.E.; Kling, P.; Erkell, L.J.; Kille, P.
R;Olsson, P.E.; Kling, P.; Erkell, L.J.; Kille, P.
A;Itle: Structural and functional analysis of the rainbow trout (Oncorhyncus mykiss)
A;Atle: Structural and functional analysis of the rainbow trout (Oncorhyncus mykiss)
A;Accession: 151049
A;Accession: 151049
A;Accession: I51049
A;Accession: I510

ö Gaps ö 0; Indels Length 4; Query Match 75.0%; Score 9; DB 2; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches

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ო RESULT S43959

Gaps

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cadmium-binding pentapeptide - downy thornapple
C; Species: Datura innoxia (downy thornapple)
C; Species: Datura innoxia (downy thornapple)
C; Date: 21-May-1990 #sequence_revision 21-May-1990 #text_change 18-Jun-1993
C; Accession: A33882
R; Jackson, P.J.; Unkefer, C.J.; Doolen, J.A.; Watt, K.; Robinson, N.J.
Proc. Natl. Acad. Sci. U.S.A. 84, 6619-6623, 1987
A; Title: Poly(gamma-glutamylcysteinyl)glycine: its role in cadmium resistance in plam.
A; Reference number: A94182; MUID:88016144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      actin I - malaria parasite (Plasmodium falciparum) (fragments)
C;Species: Plasmodium falciparum
C;Species: Plasmodium falciparum
C;Species: 03-Jun-1993 #sequence_revision 28-Oct-1994 #text_change 09-Jun-2000
C;Accession: B45525
R;Wesseling, J.G.; Snijders, P.J.F.; van Someren, P.; Jansen, J.; Smits, M.A.; Schoen Mol. Blochem. Parasitol. 35, 167-176, 1989
A;Title: Stage specific expression and genomic organization of the actin genes of the A;Reference number: A45525; MUID:89364996
R-phycoerythrin gamma-A chain - red alga (Gastroclonium coulteri) (fragment) C; Species: Gastroclonium coulteri C; Dacte: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993 C; Accession: F22565 R; Klotz, A.V.; Glazer, A.N. 1985 A; Mill: Characterization of the bilin attachment sites in R-phycoerythrin. A; Reference number: A2556; MulD:85182601 A; Molecule type: protein A; Residues: 15 < KLO.
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A) Cross-references: GB:J03988
A) Note: the authors translated the codon GAA for residue 3 as C; Comment: The actin' gene contains no introns.
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llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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100.0%; Pred. No. 0;
Live 0; Mismatches
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100.0%; Pred. No. 0;
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A; Status: preliminary
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C;Species: Triatcoma pallidipennis (assassin bug)
C;Species: Triatcoma pallidipennis (assassin bug)
C;Accession: S55238
R;Haendler, B.; Becker, A.; Noeske-Jungblut, C.; Kraetzschmar, J.; Donner, P.; Schleunin Bicchem. J. 307, 465-470, 1995
A;Title: Expression of active recombinant pallidipin, a novel platelet aggregation inhib A;Reference number: S55238; MUID:95251610
A;Accession: S55238
A;Accession: S55238
A;Accession: S55238
A;Reference number: A; S55238
A;Accession: S55238
A;Reference number: S55238
A;Reference number: S55238
A;Reference number: S55238
Ig mu chain V region (clone 13) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 20-Oct-1994 #sequence_revision 26-Jul-1996 #text_change 17-Mar-1999
C;Accession: 843959
R;Wagner, S.D.; Williams, G.T.; Larson, T.; Neuberger, M.S.; Kitamura, D.; Rajewsky, K.; Wolcleic Acids Res. 22, 1389-1393, 1994
A;Title: Antibodies generated from human immunoglobulin miniloci in transgenic mice. A;Reference number: 843956; MuID:94248036
A;Accession: 843959
A;Accession: 843956
A;Molecule type: DNA
A;Residues: 1-4 <WAG>
C;Keywords: immunoglobulin
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R-phycoerythrin alpha-2 chain - red alga (Gastroclonium coulteri) (fragment)
C;Species: Gastroclonium coulteri
C;Species: Gastroclonium coulteri
C;Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
C;Accession: B22565
R;Klotz, A.V.; Glazer, A.N.
J. Biol. Chem. 260, 4856-4863, 1985
A;Title: Characterization of the bilin attachment sites in R-phycoerythrin. A;Reference number: A22565; MUID:85182601
A;Accession: B22565
A;Molecule type: protein
A;Residues: 1-5 <KLO>
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llarity 100.0%; Pred. No. 0;
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llarity 100.0%; Pred. No. 0;
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T protein - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 12-Aug-1996 #Sequence_revision 12-Aug-1996 #text_change 05-Nov-1999
C;Accession: 137263
R;Waeber, G.; Habener, J.F.
R;Waeber, G.; Habener, J.F.
A;Title: Novel testis germ cell:specific transcript of the CREB gene contains an alte.
A;Reference number: 137263; MUID:93010691
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C. Species: Canis lupus familiaris (dog)

C. Species: Devar 1997

C. Species: S29881

S. Mulchaug, M.O.; Post, R.L.; Saccomani, G.; Leonard, R.T.; Briskin, D.P.

J. Biol. Chem. 260, 3852-3859, 1985

A. Title: Structural relatedness of three ion-transport adenosine triphosphatases arou A. Accession: S29881; MUID: 85131201
                                                                                                                                                                                                                           protamine Pl - gorilla (fragment)
C;Species: Gorilla gorilla (gorilla)
C;Species: Gorilla gorilla (gorilla)
C;Species: Gorilla gorilla)
C;Species: Gorilla gorilla (gorilla)
C;Species: 04-001-1996 #sequence_revision 04-0ct-1996 #text_change 21-Jul-2000
C;Accession: 137027
R;Queralt, R.; Oliva, R.
Gene 137, 197-204, 1993
A;Title: Identification of conserved potential regulatory sequences of the protamine-A;Reference number: 137013; MUID:94040810
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C; Genetics:
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A;Molecule type: DNA
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100.0%; Pred. No. 0;
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                                                                                hemoglobin, extracellular, chain b - earthworm (Lumbricus terrestris) (fragment)
C;Species: Lumbricus terrestris (common earthworm)
C;Species: Lumbricus terrestris (common earthworm)
C;Date: 06-Dec-1996 #sequence_revision 13-Mar-1997
C;Accession: S65726
R;Fushitani, K.; Higashiyama, K.; Asao, M.; Hosokawa, K.
Biochim. Biophys. Acta 1292, 273-280, 1996
A;Title: Characterization of the constituent polypeptides of the extracellular hemoglobine recent mumber: S65721; MUID:96176855
A;Accession: S65726
A;Accession: S65726
A;Accession S65727
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A;Ac
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C;Species: Streptomyces willmorei
C;Species: Streptomyces willmorei
C;Species: Streptomyces willmorei
C;Species: Streptomyces willmorei
C;Accession: J00355
R;Tsuda, Y:; Okada, Y:; Tanaka, M.; Shigematsu, N.; Hori, Y.; Goto, T.; Hashimoto, M.
C;Accession: J00355
R;Tsuda, Y:; Okada, Y:; Tanaka, M.; Shigematsu, N.; Hori, Y.; Goto, T.; Hashimoto, M.
R;Tsuda, Y:; Okada, Y:; Tanaka, M.; Shigematsu, N.; Hori, Y.; Goto, T.; Hashimoto, M.
A;Title: Structure and synthesis of an immunoactive lipopeptide, WS1279, of microbial
A;Reference number: J00355; MUID:91300586
A;Reference number: J00355
A;Molecule type: protein
A;Residues: 1-6 cTSU
A;Residues: 1-6 cTSU
C;Reywords: blocked amino end; lipoprotein
F;L/Binding site: sn-2,3-diacy1glycerol (Cys) (covalent) #status experimental
F;L/Modified site: fatty acylated amino end (Cys) #status experimental
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C22565
C.Paptocerythrin beta-1 chain - red alga (Gastroclonium coulteri) (fragment)
C.Species: Gastroclonium coulteri
C.Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
C.Accession: C22565
C.Accession: C22565
A.Title: Characterization of the bilin attachment sites in R-phycoerythrin. A;Reference number: A22565
A.Title: Characterization of the bilin attachment sites in R-phycoerythrin. A;Reference number: A22565
A.Molecule type: protein
A;Residues: 1-6 < KLO>
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C.Species: Box primigenius taurus (cattle)
C.Species: Box primigenius taurus (cattle)
C.Species: Box primigenius taurus (N.S. Biochem: Mol. Biol. Int. 29, 545-554, 1993
B.Chem. Mol. Biol. Int. 29, 545-554, 1993
A.Title: The major fat-globule membrane proteins, bovine components 15/16 and guinea-pig
A.Title: The major fat-globule membrane proteins, bovine components 15/16 and guinea-pig
A.Title: The major fat-globule membrane proteins, bovine components 15/16 and guinea-pig
A.Title: The major fat-globule membrane proteins
A.Status: H8394
A.Status: preliminary
A.Molecule type: protein
A.Status: A.Status: milk
A.Status: A.Status: milk
A.Status: acquence extracted from NCBI backbone (NCBIP:131518)
C; Keywords: ATP; heterodimer; hydrolase; ion transport; osmoregulation; phosphoprotein; F;4/Active site: Asp (aspartylphosphate intermediate) *status predicted
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Copyright (c) 1993 - 2000 Compugen Ltd.
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                     Lote C.J., Weiss J.B.;
"Identification in urine of a low-molecular-weight highly polar
"identification in urine of a low-molecular-weight highly polar
glycopeptide containing cysteinyl-galactose.";
Biochem. J. 123:25P-25P(1971).
--- FUNCTION: THE IDENTITY OF THE GLYCOPROTEIN FROM WHICH THIS PEPTIDE
IS DERIVED IS UNKNOWN. NO PHYSIOLOGICAL FUNCTION HAS BEEN
ATTRIBUTED. AN EXPTHROUYTE MEMBRANE GLYCOPEPTIDE HAVING A
SIMILAR STRUCTURE HAS ALSO BEEN FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Isolation and identification of a cardioactive peptide from Tenebrio
                                                                                                                                                                                                                      Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 SPECIES=C.MAENAS; TISSUE-PERICARDIAL ORGANS;
Stangier J., Hilbich C., Beyreuther K., Keller R.;
"Unusual cardioactive peptide (CCAP) from pericardial organs of the
Shore crab Carcinus maenas.";
Proc. Natl. Acad. Sci. U.S.A. 84:575-579(1987).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SPECIES=M.SEXTA;
MEDLINE=93050243; PubMed=1426284;
Cheung C.C., Loi P.K., Sylwester A.W., Lee T.D., Tublitz N.J.;
Primary structure of a cardioactive neuropeptide from the tobacco hawkmoth, Manduca sexta.";
PEBS Lett. 313:165-168(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  molitor and Spodoptera eridania.";
Bol. Chem. Hoppe-Seyler 374:1065-1074(1993).
-!- FUNCTION: THE EFFECT OF CCAP IS BOTH INO- AND CHRONOTROPIC.
-!- TISSUE SPECIFICITY: STORED IN PERICARDIAL ORGANS AND RELEASED INTO THE HEMOLYMPH.
PIN: A26363; A3633.
PIR; S27233; S27233.
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                                                                                                                                                                                                                                                                                                                                                                                                                   Manduca sexta (Tobacco hawkmoth) (Tobacco hornworm),
Tenebrio molitor (Yellow mealworm), and
Spodoptera eridania (Southern armyworm).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
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MEDLINE-94176032; PubMed-8129851;
Furuya K., Liao S., Reynolds S.E., Ota R.B., Hackett M.,
Schooley D.A.;
                                                                                                                                                                                              Length 8;
                                                                                                                                               1 1 S-LINKED (GAL. . .).
8 AA; 855 MW; C2D87AA1F5B1EB1E CRC64;
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C5A861A9CDD44EB9 CRC64;
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01-OCT-1994 (Rel. 30, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
CARDIOACTIVE PEPTIDE (CCAR).
CARCIONS MAENAS (COMMON Shore crab) (Green crab),
                                                                                                                                                                                            75.0%; Score 9; DB 1;
llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                9 AA.
                                                                                                                                                                                                                                                                                                                                  PRT;
            MEDLINE=72062338; PubMed=5126885;
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                                                                                                                                                                                                                                                                                                                                  STANDARD;
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                   Glycoprotein.
                                                                                                                                                                                                                                                                                                                                CCAP_CARMA
P38556;
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SEQUENCE
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SEQUENCE
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SEQUENCE.
MEDULNE-88058932; PubMed-3680228;
CTUZ L.7., de Santos V., Zafaralla G.C., Ramilo C.A., Zeikus R.D.,
Gray W.R., Olivera B.M.;
Gray W.R., Olivera B.M.;
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MEDILINE-88058932; PubMed-3680228;
Cruz LJ., de Santos V., Zafaralla G.C., Ramilo C.A., Zeikus R.D.,
Gray W.R., Olivera B.M.;
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Gray W.R., Olivera B.M., Cruz L.J.;
Peptide toxins from venomous Conus snails.";
Annu. Rev. Biochem. 57:665-700(1988).
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conus striatus (Striated cone).
Eukaryota; Metazoa; Mollusca; Gastropoda; Caenogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
                                                                                                                                                                                                                                                                                                                                                                  Gastropoda; Caenogastropoda;
                                                      0; Indels
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              Length 9;
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D4FC276EB4540059 CRC64;
                                                                                                                                                                                                                                                                  01-NOV-1988 (Rel. 09, Created)
01-NOV-1988 (Rel. 09, Last sequence update)
10-NOV-1995 (Rel. 32, Last annotation update)
LYS-CONOPRESSIN:G.
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01-NOV-1988 (Rel. 09, Last sequence update)
101-NOV-1995 (Rel. 32, Last annotation update)
ARG-CONOPRESSIN'S.
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Neogastropoda; Conoidea; Conidae; Conus.
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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Pred. No.
              Score 9; I
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75.0%; Scc.
100.0%; Pre
0;
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100.0%; Pr
tive 0;
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9
1037 MW;
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            Query Match
Best Local Similarity
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Best Local Similarity
Matches 1; Conserv
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P05487;
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SEQUENCE
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REVIEW

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TISSUE-PITUITARY.

Acher R., Chauvet J., Chauvet M.-T., Crepy D.;

Acher Carp. (Cyprinus carpio). Comparison with hormones from sea water bony fishs.";

Comp. abothem. "physiol. 14:245-254(1965).

-! SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.

PINIARPRO; IPROMO981;

INTERPRO; IPROMO981;

PRAM; PF00220; hormone4; 1.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rydberg B., Hall J., Karran P.;
"Active site amino acid sequence of the bovine O6-methylguanine-DNA methyltransferase."

Nucleic Acids Res. 18:17-21(1990)

-!- FUNCTION: REPAIR OF ALKYLGATED GUANINE IN DNA BY STOICHIOMETRICALLY TRANSFERRING THE ALKYL GROUP AT THE O-6 POSITION TO A CYSTEINE RESIDUE IN THE ENZYME. THIS IS A SUICIDE REACTION: THE ENZYME IS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
01-DEC-1996 (Rel. 34, Last annotation update)
METHYLATED-DNA--PROTEIN-CYSTEINE METHYLTRANSFERASE (EC 2.1.1.63) (6-0-
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INTERPRO; IPR001497; -.
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CATALYTIC ACTIVITY: DNA (CONTAINING O6-METHYLGUANINE) + PROTEIN
L-CYSTEINE = DNA (WITHOUT O6-METHYLGUANINE) + PROTEIN S-METHYL-
L-CYSTEINE
                                                                                                                                                                                                                                      Eŭkaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Actinopterygii; Neopterygii, Teleostei, Euteleostei, Ostariophysi,
Cypriniformes, Cyprinidae, Cyprininae, Cyprinus.
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
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                                                                                                    01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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100.0%; Pred. No. v.
                                                      9 AA.
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                                                                                                                                                                                                                Cyprinus carpio (Common carp).
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                                                   STANDARD;
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DISULFID 1
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Best Local Similarity
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TISSUE-THYMUS;
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P29177;
                                                   ISOT_CYPCA
P42993;
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SEQUENCE
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                             SOT_CYPCA
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MEDLINE-8807707; PubMed=3689410;
Proux J.P., Miller C.A., Li J.P., Carney R.L., Girardie A.,
Delaage M., Schooley D.A.;
"Identification of an arginine vasopressin-like diuretic hormone from
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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01-AUG-1990 (Rel. 15, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
LOCUPRESSIN (DIURETIC NEUROPEPPIDE F1/F2).
LOCUPRESTIC WINDERTORY locust).
Locusta migratoria (Migratory locust).
Eukaryota; Metazoa; Arthropoda; Trachostata; Hexapoda; Insecta; Petrygota; Neoptera; Orthopteroidea; Orthoptera; Orthoptera; Acridomorpha; Acridoidea; Acrididae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Locusta migratoria.";
Biochem. Biochem.
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INTERCHAIN (WITH C-6') (IN F2).
INTERCHAIN (WITH C-1') (IN F2).
AMIDATION.
56EB176EB451A057 CRC64;
peptides from Conus geographus and Conus straitus venoms."; J. Biol. Chem. 262:15821-15824(1987).
                                                                                                                       Gray W.R., Olivera B.M., Cruz L.J.,
Peptide toxins from venomous Conus snails.";
Annu. Rev. Biochem. 57:665-700(1988).
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; B28495; B28495
INTERPRO, IPRO00981; ...
PRAM: PF00220; hormone4; 1.
PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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17EB176EB4540050 CRC64;
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o. 0;
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larity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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llarity 100.0%; Pred. No.
Conservative 0; Mismatch
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                                                                                                       MEDLINE=89024586; PubMed=3052286;
                                                                                                                                                                                                                                                                                                                                                                                                                         9 AA; 1031 MW;
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Best Local Similarity
Matches 1; Conserv
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Matches 1; Conserv
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SEQUENCE
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Matches 1; Conserv
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OXYF_SCYCA
ID OXYF_SCYCA
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SEQUENCE
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Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyllorhinidae; Scyllorhinus.
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Squalus acanthias (Spiny dogfish).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchil; Squalea; Squaloidel; Squalidae; Squalus.
                                                        ALKYL GROUP ACCEPTOR (BY SIMILARITY).
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MEDLINE-95062247; PubMed-7972045;
MEDLINE-95062247; PubMed-7972045;
MEDLINE-95062247; PubMed-7972045;
MEDLINE-95062247; PubMed-7972045;
"Special evolution of neurohypophysial hormones in cartilaginous fishes: asvatocin and phasvatocin, two oxytocin-like peptides isolated from the spotted dogfish (Scyliorhinus caniculus).";
Proc. Natl. Acad. Sci. U.S.A. 91:11266-11270(1994).
-: FUNCTION: DISPLAYS OXYTOCIC ACTIVITY ON RAT UTERUS.
-: SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
                                                                                                                                                     75.0%; Score 9; DB 1;
100.0%; Pred. No. 0;
iive 0; Mismatches
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100.0%; Pred. No. 0;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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                  DNA repair; Transferase; Methyltransferase.
NON_TER 1 1
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Acher R., Chauvet J., Chauvet M.-T.;
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PROSITE; PS00374; MGMT; PARTIAL
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                                                                                            967 MW;
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Best Local Similarity
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Best Local Similarity
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                                                                                              9 AA;
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P42999;
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"Identification of 2 new neurohypophyseal hormones, valitocin (Val8-oxytocin) and aspartocin (Asn4-oxytocin) in a selachian fish, the spiny dog-fish (Squalus acanthias).":

C. R. Acad. Sci., D. Sci. Nat. 274:313-316(1972).

-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
INTERPRO: IPRO00981;
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PFAM: PF00220; hormone4; 1.

PROSTIE: PS00264; NEUROHYPOPHYS_HORM; 1.
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Bukaryota, Metazoa, Chordata, Cranlata; Vertebrata, Chondrichthyes,
Elasmobranchii, Galeomorphii, Galeoidea, Carcharhiniformes;
Scyllorhinidae; Scyllorhinus.
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MEDLINE-95062247; PubMed=7972045;

MEDLINE-95062247; PubMed=7972045;

MEDLINE-95062247; PubMed=7972045;

"Special evolution of neurohypophysial hormones in cartilaginous fishes: asvatocin and phasvatocin, two oxytocin-like peptides isolated from the spotted dogfish (Scyliorhinus caniculus).";

Proc. Natl. Acad. Sci. U.S.A. 91:11266-11270(1994).

-!- FUNCTION: DISPLAYS OXYTOCIC ACTIVITY ON RAT UTERUS.

-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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"Phylogeny of the neurohypophysial hormones. Two new ac
isolated from a cartilaginous fish, Squalus acanthias."
Eur. J. Biochem. 29:12-19(1972).
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5. 0;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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100.0%; Pre
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100.0%; Pro
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9 AA; 1016 MW;
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Gaps

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Indels

9 AA.

Length 9;

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"Annetocin: an oxytocin-related peptide isolated from the earthworm, Eisenia foetida:";
Bisenia foetida:";
Bischia foetida:";
Bischia foetida:";
Bischia foetida:";
Bischia Bischia: Bischia B
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Eukaryota; Metazoa; Annelida; Clitellata; Oligochaeta; Haplotaxida;
Lumbricina; Lumbricidae; Eisenia.
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Incirrata; Octopodidae; Octopus.
                                                                                    AMIDATION.
17EB176EB456D04B CRC64;
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9 AA; 996 MW; D4EEB76EB45412C9 CRC64;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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(Rel. 21, Last sequence update)
(Rel. 32, Last annotation update)
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100.0%; Pred. No. 0;
Live 0; Mismatches
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MEDLINE-94121660; PubMed-8292046;
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100.0%; Pre-
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                                                                                                                          9 AA; 1053 MW;
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TISSUE-NERVE ENDINGS;
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Matches 1; Conserv
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P80027;
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OXYT_OCTVU
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DT 01-NAR.
DC CEPHALK
OC CENARY
OC INCIPERY
OC INCIPERY
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Acher R., Chauvet J., Chauvet M.-T., Crepy D.;
"Characterization of neurohypophyseal hormones from a fresh water bony
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        fish, the carp (Cyprinus carpio). Comparison with hormones from sea water bony fishs.";
OXYI_BUFRE STANDARD; PRT; 9 AA.
P42995;
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
SERITOCIN:
SERITOCIN:
Substrace (African toad).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Bufonidae;
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Lane T.F., Sower S.A., Kawauchi H.;
Lane T.F., Sower S.A., Kawauchi H.;
Arginine vasotoin from the pituitary gland of the lamprey (Petromyzon marinus): isolation and amino acid sequence.";
Gen. Comp. Endocrinol. 70:152-157(1988).
-: FUNCTION: ANTIDIUREFIC HORMONE.
-: SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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9 AA; 983 MW; 17FF476EA5A6D04B CRC64;
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100.0%; Pred. No. 0;
iive 0; Mismatches
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01-NOV-1991 (Rel. 20, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
VASCYCCIN.
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PFAM; PF00220; hormone4; 1.
PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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PIR; S06375; S06375.
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OXYI_CYPCA
AC P23879;
DT 01-NOV-1991
DT 01-NOV-1991
DT 01-NOV-1991
DE VASOTOCIN.
OC CYPTINUS CAT
OC ACTINOPTERY
OC ACTINOPTERY
OC ACTINOPTERY
CYPTINITORNE
RP SEQUENCE.
RC SPECIES—C.CA
RA ACHER N. CA
RT GEN, CHE CA
RT GEN, CHE
CC - ! FUNCTION

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0; Indels

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                               There is now peptide of the oxytocin/vasopressin family isolated from nerves of the cephalopod Octopus vulgaris.";

Neurosci. Lett. 134:191-194(1992).

C. -!- FUNCTION: HAS A ROLE IN THE NEUROSECRETORY SYSTEM OF THE VENA C. CAVA.

INTERPRO. IPRO00981.

R PRAM: PF00220; hormone4; 1.

R PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.

R HOLDORS; AMIDATION.

T MOD_RES 9 AA: 1072 MW; 17FF476EB45409DB CRC64;
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MEDLINE=92270139; PubMed=1589145;
Reich G.;
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Search completed: February 5, 2001, 12:06:02 Job time: 279 sec

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Q91098 manorina me Q47063 escherichia Q15999 homo sapien Q99887 homo sapien Q94405 macropus ru Q94406 macropus ru Q94400 monodelphis Q38340 lactococcus Q95438 prattus norv Q69473 human herpe C12096 caprine art C12098 caprine art C12100 caprine art C12104 caprine art C12105 mycobacteri Q47475 escherichia Q48469 klebsiella Q98343 escherichia Q98439 klebsiella Q98343 homo sapien Q910318 homo sapien

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09719 human adeno
04254 fugu rubrip
02560 escherichia
012500 homo sapien
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                  GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
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ALIGNMENTS

Q47475 Q48469 Q9K343 Q9UVW2

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MEDLINE=93010691; PubMed=1396344;
Waeber G., Habener J.F.;
"Novel testis germ cell-specific transcript of the CREB gene contains an alternatively spliced exon with multiple in-frame stop codons.";
Endocrinology 131:2010-2015(1992).
EMBL; X68994; CAA48780.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                 Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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SEQUENCE FROM N.A.
MEDLINE-81001892; PubMed-7407922;
MEDLINE-81001892; PubMed-7407922;
Dhar R., Chanock R.M., Lai C.J.;
"Nonviral oligonuclectides at the 5' terminus of cytoplasmic influenza viral mRNA deduced from cloned complete genomic sequences.";
Cell 21:495-500(1980).
EMBL; M25045; AAA43202.1;
                                                                                                                                                                                                                                                                                                Viruses; ssRNA positive-strand viruses, no DNA stage; Comoviridae;
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Long, nearly identical untranslated sequences at the 3' terminal
regions of the genomic RNAs of cherry leafroll virus (walnut
strain).";
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01-NOV-1998 (Tremblrel. 08, Last annotation update)
(UDORN/72) HEMAGGLUTININ (SEG 4) CDNA, 3' END (FRAGMENT).
Influenza A virus.
Viruses; SSRNA negative-strand viruses; Orthomyxoviridae;
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
C-TERMINUS OF THE VIRAL REPLICASE (FRAGMENT).
cherry leaf roll virus.
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EMBL; Z34265; CAA84019.1; -.
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                                                                                                                                                                                                                                                                                                                   MEDLINE-96427318; PubMed-8830682;
Rehm B.H.A., Ertesvag H., Valla S.;
"A new Azotobacter vinelandii mannuronan C-5-epimerase gene (algG) is part of an alg gene cluster physically organized in a manner similar to that in Pseudomonas aeruqinosa.";
J. Bacteriol. 178:5884-5889(1996).
EMBL; X87973; CAA61230.1; -.
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Bukaryota, Wetazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
NCBI_TaxID=10116;
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                                                                   Azotobacter vinelandii.
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MEDLINE=96198747; PubMed=8612486;
Yoshikawa T., Makino S., Gao X.M., Xing G.Q., Chuang D.M.,
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08, Last annotation update)
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SEQUENCE 7 AA; 663 MW; 6DDAA8787EB05350 CRC64;
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                                                                                                                                                                                                                                                           SEQUENCE FROM N.A.
                                                                                                                                                                                NCBI_TaxID=354;
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O32560 PRELIMINARY;
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01-JAN-1998 (TrEMBLrel. 05,
01-MAY-1999 (TEMBLEEL 10, 01-MAY-1999 (TEMBLEEL 10, 01-MAY-1999 (TEMBLEEL 10, PVI CORE PROTEIN (FRAGMENT)
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Best Local Similarity
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Best Local Similarity
Matches 1; Conserv
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SEQUENCE FROM N.A.
                                                         Viruses; dsDNA vi
NCBI_TaxID=28280;
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                                                Human adenovirus
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                                                                                                     Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
NCBI_TaxID=10519;
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NCBI_TaxID=85755;
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                                                                                                                                             STRAIN=GOMEN,
Crawford-Miksza L.K., Nang R.N., Schnurr D.P.;
"Molecular surveillance of strain variation in adenoviruses
acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL, AF065065; AAD03662.1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                        SEQUENCE FROM N.A.
STRAIN=KN T96-0620, S-1058, CL 68578;
Crawford-Miksza L.K., Nang R.N., Schnurr D.P.;
Crawford-Miksza L.K., Nang R.N., Schnurr D.P.;
Molecular surveillance of strain variation in adenoviruses acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065069; AAD03664.1;
EMBL; AF065067; AAD03664.1;
EMBL; AF065067; AAD03666.1;
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Last annotation update)
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Last annotation update)
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01-MAY-1999 (TrEMBLrel. 10,
01-MAY-1999 (TrEMBLrel. 10,
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                             PRELIMINARY;
                                                                                              Human adenovirus type 7.
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Best Local Similarity
Matches 1; Conserv
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                                                                                                                                     SEQUENCE FROM N.A.
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Q9YVE3;
01-MAY-1999
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Q9YVE3
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Q9YIRO
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"Alternative splicing of the sodium channel SCN8A predicts a truncated two-domain protein in fetal brain and non-neuronal cells.";
J. Biol. Chem. 272:14008-24015(1997).
EMBL; U97673; AAB80916.1;
Tonic channel.
SEQUENCE 7 AA; 730 MW; 75B72EA2C73772A0 CRC64;
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CTAWFORD-MIKSZA D.K., Nang R.N., Schnurr D.P.;
Mollecular surveillance of Errain variation in adenoviruses causing acute respiratory disease, Av 4 and Av 7a.";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL, AF065064; AAD03659.1;
EMBL, AF065062; AAD03659.1;
EMBL, AF065063; AAD03656.1;
                                                                                                                       us type 4.
viruses, no RNA stage; Adenoviridae; Mastadenovirus.
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Actinopterygii, Neopterygii, Teleostei, Euteleostei, Neoteleostei,
Acanthomorpha, Acanthopterygii, Percomorpha, Tetraodontiformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
TRUNCATED VOLTAGE-GATED SODIUM CHANNEL ALPHA SUBUNIT (FRAGMENT).
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(TrEMBLrel. 10, Created)
(TrEMBLrel. 10, Last sequence update)
(TrEMBLrel. 10, Last annotation update)
                                                                                                                                                                                                                                                                                                                                                                                                                                                               7 AA; 980 MW; 7B5EA414140322A0 CRC64;
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SEQUENCE FROM N.A.
MEDLINE=97442476; PubMed=9295353;
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PRELIMINARY;
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Best Local Similarity
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Matches 1; Conserv
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SEQUENCE FROM N.A.
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TISSUE=PLACENTA;
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                                                                                                                              TISSUE-PLACENTA;
                                                                           NCBI_TaxID=9606;
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SEQUENCE
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Q15900;
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Q9Y4X6;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       015888;
01-NOV-1996 (TERBLEEL: 01, Created)
01-NOV-1996 (TERBLEEL: 01, Last sequence update)
01-NOV-1998 (TERBLEEL: 07, Last annotation update)
01-AUG-1998 (TERBLEEL: 07, Last annotation update)
(CLONE XP15184A) (FRAGWENT).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                        Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
Escherichia.
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Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
EMBL, AJ001620; CAA04875.1; -.
                                                                                                                                                                                                   Hesslinger C., Sawers G.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases
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Last sequence update)
Last annotation update)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-AUG-1998 (TrEMBLrel. 07, Last annotation update)
PROPIONATE KINASE (FRAGMENT).
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100.0%; Pred. No. 0;
iive 0; Mismatches
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tive 0; Mismatches
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TISSUE=PLACENTA;
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Matches 1; Conserv
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                                                                         Escherichia coli
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 01-NOV-1996 (TrEWBLrel. 01, Created)
01-NOV-1996 (TrEWBLrel. 01, Last sequence update)
01-NOV-1998 (TrEWBLrel. 07, Last annotation update)
(CLONE XP7B11A). (FRAGMENT).
HOMO Sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
(CLONE XP19G12A) (FRAGMENT).
Homo sapiens (Human).
Eukaryota; Metazona; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae, Homo.
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
NUCLEAR LIM INTERACTOR (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                        975 MW; 605EA6C5BEA5A2D3 CRC64;
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100.0%; Pred. No. 0;
ative 0; Mismatches
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100.0%; Pred. No. 0;
tive 0; Mismatches
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EMBL: L32079; AAA73890.1; -.
NON_TER 1
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Hum. Mol. Genet. 0:0-0(0).
EMBL; L32083; AAA73880.1;
NON_TER 1
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                                                                                                                        SEQUENCE FROM N.A.

Drechsler M., Schumacher V., Friedrich S., Wildhardt G., Giesler S., Schumacher V., Friedrich S., Wildhardt G., Giesler S., Schnoth A., Bodem J., Royer-Pokora B., Genoric structure, alternative transcripts and chromosomal localization of the human LIM domain binding protein 1 gene LDB1/NLI.";
Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
BNBL; AJ243097; CAB45408.1; -.
NON_TER 8
SEQUENCE 8 AA; 767 MW; EEGEBDDEB862D5B6 CRC64;
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
PRO ALPHA 1 TYPE III COLLAGEN PROTEIN (FRAGMENT).
Oryctolagus cuniculus (Rabbit).
Eukaryota, Metazoa; Chordata, Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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MEDLINE-96377339; Pubmed-8783186;
Metsaranta M., Kujala U.M., Pelliniemi L., Osterman H., Aho H.,
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NCBI_TaxID=9606;
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Search completed: February 5, 2001, 12:05:38 Job time: 289 sec

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Length 8;

75.0%; Score 9; DB 6; 100.0%; Pred. No. 0; Live 0; Mismatches

Query Match 75.0 Best Local Similarity 100. Matches 1; Conservative

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RESULT
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                            GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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Maximum Match 100%
Listing first 45 summaries
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Gapop 10.0 , Gapext 0.5
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Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	Sequence which cor	Binding receptor w .	Synthetic peptide	Metal trapping pep	Metal trapping pep	IGF-1 analogue N-t	IGF-1 analogue N-t	IGF-1 analogue N-t	Human interleukin-	Anti-inflammatory	Anti-inflammatory	Anti-inflammatory
SUMMARIES	ΙD	P10301	P90982	P91660	R34022	R34023	R51439	R51440	R51441	W19847	W56231	W56211	W56200
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WPI; 1981-53498D/30 (53498D).

Anti-inflammatory Peptide used in ti. Biostatin TY252 sy Sequence of antiin Sequence of peptid Enkephalin-like an Analgesic and neur Sequence of chromo Sequence of chromo Sequence of cyclic Sequence of cyclic Sequence of cyclic Sequence of cyclic Sequence b Synthetic peptide Peptide fragment # Peptide fragment #	Thioredoxin active Cyclic platelet ag Farnesyl-protein t
3 19 W56172 3 20 W88037 3 21 P10138 4 2 P10033 4 2 P10093 4 2 P10382 4 4 P30083 4 4 P30083 4 5 P40538 4 5 P40538 4 6 P40538 4 8 P70833 4 10 P91661 4 12 R11077	4 12 R11508 4 12 R11744 4 12 R11744 4 12 R13745 4 12 R14723 4 12 R15751 4 12 R15753 4 12 R15754 4 12 R15754 4 12 R15756 4 12 R15756 4 12 R15756 4 12 R15756 4 12 R15759 4 12 R15759
27 27 27 27 27 27 27 27 27 27 27 27 27 2	
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ALIGNMENTS

Cystine peptide; insulin activity; hormone; diabetes therapy; __note= "bonded to Cys(5) of P10115, which is residue 19 of the B chain fragment B15-27" Sequence which corresp. to resides 19-21 of insulin A chain. Location/Qualifiers 2 P10301 standard; Protein; 3 AA. 30DD-0011868 79DD-0215100 19-AUG-1992 (first entry) Stange H; antidiabetic agent (LOSS/) LOSSE G. Key Disulfide-bond 01-FEB-1980; 20-AUG-1979; 29-APR-1981. DD147942-A. Losse G, P10301; Mammal

Cysteine

Sequence

Query Match

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RESULT P90982

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                                                                                                                                                                                                                                                               to residues 12-14 of naturally occurring
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        Indels
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/note="(aceto amido methyl) NH2-Cys"
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                                                                                                                                                            P91660 standard; protein; 3 AA.
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                                                                                                                                                                                                                                                             Synthetic peptide corresp. to epidermal growth factor (EGF)
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 The article for inactivating a toxic material comprises a solid carrier bearing a first receptor which binds the target ligand and a second receptor which degrades the target ligand. This synthetic peptide is a preferred first receptor. The article may be used for covering surfaces to protect or decontaminate the surface. The article is esp. for degrading toxic organic openic spp. sep. organophosphorous opds., (eg pesticides and nerve poisons, bacteria and viruses, in environmental, chemical, military and industrial settings.
                                                                                         The inventors claim the prepn. of new cysteine peptides with insulin-
                                                                                                       like activity. The new peptides comprise a sequence with corresp. to a central portion of the insulin B chain (esp. B15-27) and a cysteine-contg. peptide with an AA sequence corresp. to a terminal portion of the insulin A chain (esp. A1-7). The new cysteine peptides are antidiabetic agents with appreciable insulin activity (lower than that of insulin itself) and low antigenic activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Article for inactivating toxic materials, eg organo:phosporous cpds. comprises solid carrier bearing target ligand binding receptor and ligand-degrading receptor, pref. enzyme
                                                                                                                                                                                                                                                                                                                  Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Toxic material; inactivator; organophosphorous cpds.; nerve poison; pesticide; decontaminant; military.
peptide(s) with antidiabetic activity - contg. fragments \boldsymbol{n} A and B chains
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100.0%; Pred. No. 0;
iive 0; Mismatches
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Pred. No.
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                                                        Claim 1; Page 10; 11pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    P90982 standard; peptide; 3 AA.
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100.0%;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               AD INC
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                                                                                                                                                                                                                                                                                             Local Similarity
les 1; Conserv
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Best Local Similarity
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                      insulin
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Sequence

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/note= "OMe protected"
 of cysteine"
                                                                                                                                                                                                       (ICHI-) ICHIKAWA GOSEI KAGAKU KK.
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(GROP-) GROPEP PTY LTD.
(USDA) US SEC OF AGRIC.
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                                                                                                                                                                                                                                         WPI; 1993-021449/03.
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                 3 AA;
                 Modified-site
                                                                  JP04346999-A
                                                                                                                                     24-MAY-1991;
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R51439
ID R5
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                                                                                                                                                                                     "Cys(Acm); Acm- acetamidomethyl-protected SH of cysteine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       /note= "Cys(Acm); Acm= acetamidomethyl-protected SH
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              New acyclic peptide with metal trapping activity - exhibits in vivo metal transporting action and can be used as drug, diagnostic agent or functional material
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                                                                                Electrostatic interaction; metalloprotein; binding site;
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                                                                                                                                                                                                                                                                                        /note= "OMe protected"
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                                                 Metal trapping peptide intermediate.
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                                                                                                                                                                      Location/Qualifiers
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                 (first entry)
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Best Local Similarity
Matches 1; Conserv
                                                                                                protecting group.
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Modified-site
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                                                                                                                                     Synthetic
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The acyclic peptide (see R31340) with metal trapping activity may be produced by fragment condensation using the peptide fragments given in R34022-29.
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New acyclic peptide with metal trapping activity - exhibits in vivo metal transporting action and can be used as drug, diagnostic agent: or functional material
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in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
                                                                                                                                                                                                                                                                                                                                                                                                                               0; Indels
                                                                                                                                                                                                                                                                                                                                                                           14; Length 3;
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                                                                                                                                                                                                                                                                                                                                                                        Score 9; DB 14; Pred. No. 0; 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                    75.0%; Sco...
100.0%; Pre
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92US-0947035
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(GROP-) GROPEP PIY LID.
(USDA ) US SEC OF AGRIC.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Homo sapiens.
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Best Local Si
Matches 1;
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R51441
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IGF-1 analogues used pref. (i) lack 1-5 N-terminal amino aicds or (ii) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-1 analogue pref. comprises the N-terminal Gly in R51439-49.

IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly. Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. Comprises the N-terminal given in R51450-53.

Exp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
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in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 15; Length 3; 100.0%; Pred. No. 0; Os Indels ive 0; Mismatches 0; Indels
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            Page 37; 45pp; English
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(GROP-) GROPEP PTY LTD.
(USDA ) US SEC OF AGRIC.
                                                                                                                                                                                                                                                                N-terminal extension.
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Best Local Similarity
                                                                                                                                                                                                                                                                                           3 AA;
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                                                                                                                                                                                                                                                                                             Sequence
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             Claim
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weeks after hatch.

IGF-1 analogues used pref. (i) lack 1-5 N-terminal amino aicds or (ii) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-1 analogue pref. comprises the N-terminal given in R51439-49.

IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53.

Bsp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
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in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
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100.0%; Pred. No. 0;
ive 0; Mismatches
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Gaps

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W56171-248 represent anti-inflammatory tripeptides of the invention. They are derived from the formulae:
Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where
Xaa = any amino acid residue.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Anti-inflammatory peptides and derivatives - used for treating, e.g arthritis, ulcerative colitis, auto-immune disease, allergy asthma, shock, HIV infection, transplant rejection or Alzheimer's disease
                                                                                                                                                                                                                                                                                        Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
     Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Claim 7; Page 35; 42pp; English.
                                                                                                                                                      W56231 standard; peptide; 3 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           97US-0864301.
96US-0025376.
96US-0753141.
                                                                                                                                                                                                                                                         Anti-inflammatory tripeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              97WO-IL00295
                                                                                                                                                                                                                          20-JUL-1998 (first entry)
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     Conservative
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                              03-SEP-1997;
                                                                                                                                                                                                                                                                                                                                                                                            WO9809985-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               28-MAY-1997;
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                                                                                                                                                                                                                                                                                                                                                             Synthetic.
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W56231
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                      adjacent
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IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic: Gly and the Thr normally adjact to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53. Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid N-terminal extension.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Interleukin-12; Fc gamma-1; immunosuppressive; autoimmune disease; graft rejection; toxic shock; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               A peptide (W19847) comprises the three N-terminal amino acid residues of human interleukin-12 (IL-12) p40 subunit. It is encoded both by native human p40 cDNA (T72095) and by a primer (T72095) based on the native sequence. PCR amplification has been used to amplify p40 cDNA. A fusion protein comprising the p40 subunit and human FC gamma-1 (see also W19853) was expressed in E. coli. This fusion protein has a longer in vivo half-life than native p40 and can be used as an immunosuppressive (e.g. to treat autoimmune diseases or to inhibit graff rejection) or to treat or prevent endotoxin-induced shock.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Fusion protein containing interleukin-12 p40 sub-unit - has increased stability, used to inhibit graft rejection, or treat autoimmune disease and endotoxin-induced shock
                                                                                                                                                                                                                          ö
                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                       Length 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Human interleukin-12 p40 subunit N-terminal peptide.
                                                                                                                                                                                       DB 15;
. 0;
                                                                                                                                                                                     Score 9; DB 15
Pred. No. 0;
0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                (BETH-) BETH ISRAEL HOSPITAL ASSOC. (UYMA-) UNIV MASSACHUSETTS.
                                                                                                                                                                           75.0%; Scc.
100.0%; Pre
                                                                                                                                                                                                                                                                                                                                                                           W19847 standard; Peptide; 3 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                             (first entry)
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Best Local Similarity
'-has 1; Conserve
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N-PSDB; T72095-96.
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Best Local Similarity
                                                                                                                                        3 AA;
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                                                                                                                                        Sequence
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W19847
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Hirschberg DL;

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Cyclic derivatives of the peptides also function as anti-inflammatory agents. The peptides can be covalently linked to one another either directly or through a spacer. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus, anti-inflammatory activity. The peptides and compositions have anti-inflammatory activity. I.e. inhibitory effects against a cellular and humoral immune response, including a response not associated with inflammation. The peptides also inhibit the ability of macrophages and reals to adhere to extracellular matrix components and fibronectin, as well as up-regulated fast receptor expression in T-cells. They can be used to inhibit unwanted immune reaction and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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100.0%; Pred. No. 0;
ive 0; Mismatches
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They are derived from the formulae:

Xaa -Glu-Arg, Arg-Glu, xaa, Xaa -Arg-Glu, or Glu-arg-Xaa, where
Xaa - any amino acid residue.

Yaa any amino acid residue.

Cyclic derivatives of the peptides also function as anti-inflammatory agents. The peptides can be covalently linked to one another either directly or through a spacer. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus, anti-inflammatory activity. The peptides and compositions have anti-inflammatory activity. The peptides and compositions have anti-inmune activity, i.e. inhibitory effects against a cellular and humoral immune response, including a response not associated with inflammation. The peptides also inhibit the ability of macrophages and T-cells to adhere to extracellular matrix components and fibronectin, as well as up-regulated fas receptor expression in T-cells. They can be used to inhibit unwanted immune reaction and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                             Anti-inflammatory peptides and derivatives - used for treating, e.g. arthritis, ulcerative colitis, auto-immune disease, allergy asthma, shock, HIV infection, transplant rejection or Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W56171-248 represent anti-inflammatory tripeptides of the invention
                                            Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
                                                                                                                                                                                                                                                                                                                                               Beserman P, Eisenbachschwartz M, Hirschberg DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 5; Page 34; 42pp; English.
                                                                                                                                                                                                                                                                                                               (YEDA ) YEDA RES & DEV CO LID
             Anti-inflammatory tripeptide.
                                                                                                                                                                                                                                            97US-0864301.
96US-0025376.
96US-0753141.
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                                                                                                              Synthetic.
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Xaa-Glu-Arq, Arg-Glu-Xaa. Xaa-Arg-Glu, or Glu-arg-Xaa, where
Xaa = any amino acid residue.

Yaa-Glu-Arq, Arg-Glu-Xaa and Arg-Glu, or Glu-arg-Xaa, where
Yaa = any amino acid residue.

Cyclic derivatives of the peptides also function as anti-inflammatory
agents. The peptides can be covalently linked to one another either
directly or through a spacer. The peptides and their derivatives have
macrophage inhibitory and T-cell inhibitory activity and thus,
anti-inflammatory activity. The peptides and compositions have
anti-inmune activity, i.e. inhibitory effects against a cellular and
humoral immune response, including a response not associated with
inflammation. The peptides also inhibit the ability of macrophages and
r-cells to adhere to extracellular matrix components and fibronectin, as
well as up-regulated fas receptor expression in T-cells. They can be used
to inhibit unwanted immune reaction and inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Anti-inflammatory peptides and derivatives - used for treating, e.g. arthritis, ulcerative colitis, auto-immune disease, allergy asthma, shock, HIV infection, transplant rejection or Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              W56171-248 represent anti-inflammatory tripeptides of the invention.
                                                                                                                                                            Anti-inflammatory, macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
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100.0%; Pred. No. 0;
live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Eisenbachschwartz M,
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                                W56211 standard; peptide; 3 AA.
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                                                                                                                              Anti-inflammatory tripeptide.
                                                                                                                                                                                                                                                                                                                                                             97US-0864301.
96US-0025376.
96US-0753141.
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                                                                                             (first entry)
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                                                                                                                                                                                                                               Synthetic.
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W56200
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Gaps
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0
  DB 19; Length 3; . 0;
                                                0; Indels
75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                W56172 standard; peptide; 3 AA.
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                                                Conservative
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Solid phase synthesis; biostatin; cytostatic; tyrosine kinase inhibitor; cellular proliferation inhibition; somatostatin; antitumor.
                                                                                                                                                                                      Peptides W88033-39 are used in the hair composition of the invention. The specification describes a hair composition that comprises at least one oligoceptide having more than two cysteine residues and more
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Biostatin preparation in high yield by solid synthesis, including closure of disulfide bridge before cleavage from support, useful as antitumor agent -
                                                                                                    New composition for hair - comprises oligopeptide having more than cysteine residues and reducing agent
                                                                                                                                                                                                                                         than three amino acid residues of the same kind (except cysteine) a reducing agent. The composition is useful for styling hair to take various shapes and waves safely and effectively.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note= "Thr(tBu) with C-terminal amide group"
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Pred. No. 0;
; Mismatches 0;
                                                                                                                                                                                                                                                                                                                                                  Score 9; DB
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                                                                                                                                                      Example 2; Page 6; 17pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                  75.0%; Scc.
100.0%; Pre
0; }
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   "Cys(Acm)"
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Y51896 standard; peptide; 3 AA.
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97JP-0161657
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2000-224663/19
                                                                   WPI; 1999-148442/13
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Best Local Similarity
                                 (LIOY ) LION CORP.
                                                                                                                                                                                                                                                                                                                3 AA;
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18-JUN-1997;
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Cyclic derivatives of the peptides also function as anti-inflammatory agents. The peptides can be covalently linked to one another either directly or through a spacer. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus, anti-inflammatory activity. The peptides and compositions have anti-inflammatory activity. The peptides and compositions have incliamme activity, i.e. inhibitory effects against a cellular and humoral immune response, including a response not associated with inflammation. The peptides also inhibit the ability of macrophages and T-cells to adhere to extracellular matrix components and fibronectin, as well as up-regulated fas receptor expression in T-cells. They can be used to inhibit unwanted immune reaction and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                         Anti-inflammatory peptides and derivatives - used for treating, e.g. arthritis, ulcerative colitis, auto-immune disease, allergy asthma, shock, HIV infection, transplant rejection or Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                 W56171-248 represent anti-inflammatory tripeptides of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ö
                                                                                                                                                                                                                                                                                                                                                                                                                               They are derived from the formulae:
Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 19; Length 3; . 0;
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                                                                                                                                                                                                                                           Hirschberg
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                                                                                                                                                                                                                                         Eisenbachschwartz M,
                                                                                                                                                                                                                                                                                                                                                                                 Claim 3; Page 34; 42pp; English
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    W88037 standard; peptide; 3 AA.
                                                                                                                                                                                                         (YEDA ) YEDA RES & DEV CO LTD
                                                                                                                                    97US-0864301.
96US-0025376.
96US-0753141.
                                                                                                    97WO-IL00295
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Best Local Similarity
Matches 1; Conserv
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                               WO9809985-A2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Peptide used
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                JP11012138-A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 18-JUN-1997;
                                                                                                    03-SEP-1997;
                                                                                                                                                                        20-NOV-1996;
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                                                                 12-MAR-1998
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Synthetic
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Gaps

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Indels Length 3;

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This invention describes a novel method for the solid phase (SP)

Synthesis of biostatin (TT 232) (I) which includes closing the disulfide bridge by oxidation of the completely or partially constructed peptide while still bonded to the solid phase. The products of the invention have cytostatic activity and are tyrosine kinase inhibitors. (I) inhibits the tyrosine kinase activity of various human stomach cancer cell lines and thus inhibits cellular proliferation. The heptapeptide (I) described in the invention is a somatostatin analog which shows strong antitumor cativity in vitro and in vivo. The SP synthesis method gives (I) is more easily and in markedly higher yield than by the method of EP50580, in which the cyclization is carried out after cleaving the peptide from the casin. The solution method is also a simple synthesis of (I) in high yield of typically the tert-butyl-protected precursor can be oxidized in a yield of 70-80%. Y51896-Y51900 represent peptides used in the synthesis of biostatin TT232 described in the method of the invention.
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3 AA; Sequence

0; Gaps 0; Indels Query Match 75.0%; Score 9; DB 21; Length 3; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches 0; Indels

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Search completed: February 5, 2001, 12:01:19 Job time: 366 sec

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Sequence Sequence Sequence Sequence Sequence

Appl Appl Appl Appl

Sequence

Appl Appl Appl

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GENERAL INFORMATION:

APPLICANT: MORISHITA, Hideaki
APPLICANT: MARAMORI, Toshinori
APPLICANT: KANAMORI, Toshinori
APPLICANT: KANAMORI, Toshinori
APPLICANT: KANAMORI, Toshinori
APPLICANT: KANAMORI, TOSHINORI
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE SAME, TITLE OF INVENTION: INHIBITION PROCESS, DRUG COMPOSITION AND TITLE OF INVENTION: TREATING USING THE SAME
CORRESPONDENCE: 108
CORRESPONDENCE ADDRESS:
ADDRESSEE: Burns, Doane, Swecker & Mathis
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: BAPC compatible
SOFTWARE: PAPELICATION DATA:
APPLICATION NUMBER: US/07/791,213D
FILING DATE: 13-NOV-1991
CLASSIFICATION: 435
PRIOR APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               .;
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US-08-079-812-31
US-08-122-510-11
US-08-122-510-12
US-08-122-510-13
US-08-122-510-13
US-08-122-510-13
US-08-122-455-8
US-08-372-455-8
US-08-371-585A-1
US-08-321-585A-1
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-231-1505A-14
US-08-231-1505A-14
US-08-231-150A-28
US-08-231-150A-28
US-08-231-150A-3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Burns, Doane, Swecker & Mathis STREET: P.O. Box 1404
CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
Live 0; Mismatches
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FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                             ; Sequence 23, Application US/07791213D
; Patent No. 5409895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 36,607
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TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (703) 836-6620
TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 23:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            STATE: Virginia
COUNTRY: United States
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              NAME: Meuth, Donna M
REGISTRATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            MOLECULE TYPE: peptide
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3Y: linear
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Best Local Similarity
Matches 1; Conserv
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US-07-791-213D-23
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     23, Appl
39, Appl
9, Appli
307, App
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307, App
307, App
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9, Appli
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/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6_COMB.pep:*
/cgn2_6/ptodata/2/iaa/PcTuS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/PcTuS_COMB.pep:*
                       GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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US-08-354-240A-12

US-08-354-150A-23

US-08-293-150A-39

US-08-461-386-9

US-08-461-386-9

US-08-461-386-9

US-08-486-399-307

US-08-486-397-307

US-08-486-397-307

US-08-486-397-307

US-09-249-471-307

US-09-249-471-307

US-09-249-451-307

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US-07-941-307

US-07-941-307

US-07-941-307

US-07-941-307

US-07-941-307

US-07-941-303
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                                                                                                                                                                                                                                                                                                                        174772 segs, 17957048 residues
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                                                                                                                              2001, 11:56:18
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Maximum Match 100%
Listing first 45 summaries
                                                                                            protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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length: 2000000000
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Database

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Title: Perfect score:

Sequence:

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Searched:

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WITLE OF INVENTION: Imaging UMBER OF SEQUENCES: 11
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                                                                                                                                                COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-133-804-9
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                                                                                                                                                                                                                                                                                                POLYPEPTIDE, DNA FRAGMENT ENCODING THE SAME AND ENZYME SAME AND PROCESS FOR PRODUCING THE SAME, AND ENZYME INHIBITION PROCESS, DRUG COMPOSITION AND METHODS OF TREATING USING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSE: Burns, Doane, Swecker & Mathis STREET: P.O. Box 1404
CITY: Alexandria
STATE: Virginia
STATE: Virginia
CONNTRY: United States
ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/791,213D
FILLING DATE: 13-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
Live 0; Mismatches
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APPLICATION NUMBER: JP 2-306745
FILING DATE: 13-NOV-1990
ATTORNEY AGENT INFORMATION:
NAME: Meuth, Donna M
EGISTRATION NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 029650-032
REFERENCE/DOCKET NUMBER: 029650-032
TELECHONE: (703) 836-620
TELECHONE: (703) 836-620
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 anino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9, Application US/08133804
Patent No. 5534254
GENERAL INFORMATION:
APPLICANT: Huston, James S.
APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Blosynthetic B
                                                                                                                                                                    Sequence 39, Application US/07791213D Patent No. 5409895
                                                                                                                                                                                                          GENERAL INFORMATION:
APPLICANT: MORISHITA, Hideaki
APPLICANT: RANAMORI, TOSHIDOLI
APPLICANT: ROBUHARA, MASANITO
TITLE OF INVENTION: SAME AND PROFITLE OF INVENTION: SAME AND PROFITLE OF INVENTION: INHIBITION PROFITLE OF INVENTION: TREATING USIN NUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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Best Local Similarity
Matches 1; Conserva
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                                                                                                                            RESULT 2
US-07-791-213D-39
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US-08-133-804-9
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                                               Thibeault/Patent Department
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OTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COUNTRY: USA
ZIP: 5317-1514
COMPUTER READABLE FORM:
MEDTUM TYPE: 'FIOPPY disk
COMPUTER: IBM, PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/354,240A
FILING DATE: '12-DEC-1994
                                                                                                                                                                                                                                                      SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/133,804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Sequence 12, Application US/08354240A
Patent No. 5670356
GENERAL INFORMATION:
APPLICANT: Sherf, Bruce A.
APPLICANT: Wood, Keith V.
TITLE OF INVENTION: MODIFIED LUCIFERASE
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSE: DeWitt Ross & Stevens, S.C.
STREET: 8000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
ive 0; Mismatches
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSE: Testa, Hurwitz & Thibeault/P
STREET: Exchange Place, 53 State Street
CITY: Boston
                                                                                                                                                 ZIP: 02109
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 424
ATTORNEX/AGENT INFORMATION:
NAME: KELIEY, ROAID D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054,
TELECOMMUICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEPHONE: 617-248-7477
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              sss: single
linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 2 amino acids
                                                                                                Massachusetts
: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME/KEY: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Madison
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              STRANDEDNESS:
                                                                                                                                                                                                                                                                                                                                FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   US-08-354-240A-12
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RESULT 6
US-08-293-150A-23
Sequence 23, Application US/08293150A
Patent No. 5792629
GENERAL INFORMATION:
APPLICANT: MORIBITA, Hideaki
APPLICANT: MOBUHARA, Masahiro
TITLE OF INVENTION: POLYPEPTIDE,
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE
TITLE OF INVENTION: INHIBITION PROCESS, DRUG COMPOSITION AND METHODS OF
TITLE OF INVENTION: TREATING USING THE SAME, AND ENZYME
TITLE OF INVENTION: TREATING USING THE SAME
                                                              Gaps
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                                                              0; Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Length 2;
                       Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
STREET: P.O. Box 1404
CITY: Alexandria
STATE: Virginia
COUNTRY: United States
LIP: 22313-1404
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER READBLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OMPUTER: DAY PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/293,150A
FILING DATE: 19-AUG-1994
CLASSIFICATION: 514
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/791,213
FILING DATE: 13-NOV-1990
PRIOR APPLICATION NUMBER: 3P 2-306745
FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: MEULIN, DORNA M.
REGISTRATION NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 30,607
REFERENCE/DOCKET NUMBER: 30,607
                       DB 1;
5. 0;
          75.0%; Score ,,
100.0%; Pred. No. 0;
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US-08-293-150A-39
; Sequence 39, Application US/08293150A
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100.0%; Pre
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (703) 836-620
TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                              Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; TOPOLOGY: linear
; MOLECULE TYPE: peptide
US-08-293-150A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Ouery Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  amino acid
                       Query Match
Best Local Similarity
                                                                                                      1 C 1
                                                                                                                                          2 C 2
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                                                              Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     GENERAL INFORMATION:
APPLICANT: Huston, James S.
APPLICANT: Huston, James S.
APPLICANT: Oppermann, Hermann
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    LOCATION: 1..2

CTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"

US-08-461-838-9
                                                                                                                                                                                                                                                                                                                                                                       Indels
                                                                                                                                                                                                                                                                                                                                Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     STATE: Massachusetts
COUNTRY: USA
ZIP: 02109
COMPUTER: E1009
COMPUTER: Floppy disk
MEDIUM TYPE: Floppy disk
OCHUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                         75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
Live 0; Mismatches
                NAME: Sara, Charles S.
REGIGSTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 34506.029
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-831-2106
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/08/461,838 FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CLASSIFICATION: 424
ATTORNEY/ACENT INFORMATION:
NAME: RELIEV, ROBIN:
REFERENCE/COCKET NUMBER: 34,637
REFERENCE/COCKET NUMBER: 2054/
TELECOMMUICATION INFORMATION:
TELEPHONE: 617-248-717
TELEFAX: 617-248-710
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Sequence 9, Application US/08461838 Patent No. 5753204
ATTORNEY/AGENT INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                         TOPOLOGY: linear

MOLECULE TYPE: protein

US-08-354-240A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 1; Conserva
                                                                                                                                                                                                      TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         CITY: Boston
STATE: Massacl
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 5
US-08-461-838-9
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us-09-687-267-1.rai

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STATE: Massachusetts
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              90071
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-465-380-307
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                                    COUNTRY:
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                                                                                                                                                    POLYPEPTIDE, DNA FRAGMENT ENCODING THE SAME AND FNCCESS FOR PRODUCING THE SAME, AND ENZYME INHIBITION PROCESS, DRUG COMPOSITION AND METHODS OF TREATING USING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Sequence 9, Application US/08461386
Patent No. 5837846
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: Huston, James S.
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynchetic Binding Proteins For TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 2;
                                                                                                                                                                                                                                                                                                                                  CONTESTONDEL ADURESS.

ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS STREET: P.O. Box 1404

CITY: Alexandria
STREET: P.O. Box 1404

CITY: Alexandria
STATE: Virginia
STATE: Virginia
STATE: 10.164

CONDIRY: United States
LONDING TYPE: Floppy disk
COMPUTER REAABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TEM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/293,150A
FILING DATE: 19-AUG-1994
CLASSIFICATION NUMBER: US 07/791,213
FILING DATE: 13-NOV-1990
PRIOR APPLICATION NUMBER: US 07/791,213
FILING DATE: 13-NOV-1990
PRIOR APPLICATION NUMBER: JP 2-306745
FILING DATE: 13-NOV-1990
PRIOR APPLICATION NUMBER: 36-607
REFERENCE/DOCKET NUMBER: 36-607
REFERENCE/DOCKET NUMBER: 36-607
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        75.0%; Score 9; 100.0%; Pred. No
                                                    APPLICANT: MORISHITA, Hideaki APPLICANT: KANAMORI, Toshinori APPLICANT: NOBUHARA, Masahiro: TITLE OF INVENTION: SAME AND PROCITITE OF INVENTION: SAME AND PROCITITE OF INVENTION: INHIBITION PITITLE OF INVENTION: TREATING USIN UNMBER OF SEQUENCES: 110
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACIERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  LENGIH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           MOLECULE TYPE: peptide US-08-293-150A-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Query Match
Best Local Similarity
Matches 1; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                RESULT 8
US-08-461-386-9
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ; LOCATION: 1..2
; OTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"
US-08-461-386-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: George P. Vlasuk, Patric H. Stanssens, APPLICANT: George P. Vlasuk, Patric H. Stanssens, APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys, APPLICANT: Year R. Laroche, Laurent S. Jespers, APPLICANT: Yannick G.J. Gansemans, Matthew Moyle, APPLICANT: Peter W. Bergum
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT TITLE OF INVENTION: PROTEIN
WUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
ZIP: 02109

COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOTTWARE: PatentIn Release #1.0, Version #1.25
APPLICATION NUMBER: US/08/461,386
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            75.0%; Score 9; DB 2;
100.0%; Pred. No. 0;
iive 0; Mismatches
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MEDIUM TYPE: .3.5" Diskette, 1.44 Mb
MEDIUM TYPE: .storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,380
FILING DATE: June 5, 1995
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 307, Application US/08465380 Patent No. 5863894
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                           CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: KELLEY, ROBIN D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054
TELECOMMUNICATION INFORMATION:
TELEFAN: 617-248-7477
TELEFAN: 617-248-710
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TYPE: amino acid
STRANDEDNESS: single
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Gaps

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0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                             Length 2;
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TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                     Xaa in location 2 is an amino acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      OTHER INFORMATION: Xaa in location 2 is an OTHER INFORMATION: amino acid.
                                                                                                                                                                                                                                                                                                                                                                                                                DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                         ; Pred. No. 0;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 90071
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICATION NUMBER: US/08/486,399
FILING DATE: June 5, 1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: October 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
                                                                                                                                                                                                                                                                                                                                                                                                                75.0%; Score 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 COMPUTER: IBM COMPATIBLE OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        213/270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 307, Application US/08486399; Patent No. 5865543; GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY: linear FRAGMENT TYPE: internal fragment
                                                                                                                                                          TOPOLOGY: linear
FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            REFERENCE/DOCKET NUMBER: 213/
TELECOMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                            100.08;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
         TELEX: 67-3510 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CURRENT APPLICATION DATA APPLICATION NUMBER: US
                                                                     SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                  ; OTHER INFORMATION:
; OTHER INFORMATION:
US-08-486-397-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
                                                                                                                                 amino acid
KY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                         Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 11
US-08-486-399-307
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                                                                                                                                                                                                                                                                                                                                                                                                             Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Matches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    0: Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT: George P. Vlasuk, Patric H. Stanssens, APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys, APPLICANT: Yves R. Laroche, Laurent S. Jespers, APPLICANT: Yannick G.J. Gansemans, Matthew Moyle, APPLICANT: Peter W. Bergum TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT TITLE OF INVENTION: PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FEATURE:
OTHER INFORMATION: Xaa in location 2 is an
CHER INFORMATION: amino acid.
US-08-465-380-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     DB 2;
. 0;
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100.0%; Pred. No. v.
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MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: October 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFRENCE/DOCKET NUMBER: 213/269
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFRAX: (213) 955-0440
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: 0ctober 18, 1994
ATTORNEY AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 213/268
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: (7-3510
TELEX: (7-351
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              APPLICATION NUMBER: US/08/486,397
FILING DATE: June 5, 1995
CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 307, Application US/08486397
Patent No. 5866542
GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NUMBER OF SEQUENCES: 357
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       STREET: Sulte ...
STREET: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Ouery Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                         TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                            linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: 11
FRAGMENT TYPE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: UZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-486-397-307
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US-08-486-399-307

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APPLICANT: Lauwereys, Marc Josef
APPLICANT: Laroche, Yves Rene
APPLICANT: Laroche, Yves Rene
APPLICANT: Jespers, Laurent Stephane
APPLICANT: Gansemans, Yannick Georges Jozef
APPLICANT: Moyle, Matthew
APPLICANT: Bergum, Peter W.
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
WUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
COMPUTER: 1BM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFFWARE: World Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/634,641
FILING DATE: APTIL19, 1996
PRIOR APPLICATION NUMBER: PCT/US95/13231
FILING DATE: October 17, 1995
APPLICATION NUMBER: 08/486,399
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/486,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/486,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: JUNE 5, 1995
APPLICATION NUMBER: 08/426,110
FILING DATE: SOCTOBER 18, 1994
ATTORNEY/AGERT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGIESTRATION NUMBER: 30,136
DEPENDENT OF THE NUMBER: 10,136
                                                                                                 Vlasuk, George P. Vlasuk
Stanssens, Patrick Eric Hug
Mensens, Joris Hilda Lieven
                    ; Sequence 307, Application US/08634641; Patent No. 5955294
                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         REFERENCE/DOCKET NUMBER: 21
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              SEQUENCE CHARACTERISTICS:
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OTHER INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COUNTRY: U.S.A. ZIP: 90071
                                                                            GENERAL INFORMATION:
     US-08-634-641-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             US-08-634-641-307
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                                                                                                                                                    APPLICANT:
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                                                     Gaps
                                                   0;
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                                                                                                                                                                                                                                                               Sequence 307, Application US/08461965
Patent No. 5872098
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys,
APPLICANT: Yves R. Laroche, Laurent S. Jespers,
APPLICANT: Year R. Laroche, Laurent S. Jespers,
APPLICANT: Wantlow Gensemans, Matthew Moyle,
APPLICANT: Peter W. Bergum
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon 6 Lyon
                                                   0; Indels
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     Length 2;
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  DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ); OTHER INFORMATION: Xaa in location 2 is an ; OTHER INFORMATION: amino acid. US-08-461-965-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2;
. 0;
75.0%; Score 9; DB 2
100.0%; Pred. No. 0;
tive 0; Mismatches
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100.0%; Pred. No. 0;
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WOR'D PERFECT 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 PAPLICATION NUMBER: US/08/461,965
FILING DATE: June 5, 1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: OCCODET 18, 1994
ATOMNEY GENT OCCODET 18, 1994
ATOMNEY GENT INCORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 210/243
TELECOMMUNICATION INFORMATION:
TELEDRONE: C1313, 489.1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRAGMENT TYPE: internal fragment FEATURE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Conservative
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STATE: California
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Best Local Similarity
Matches 1; Conserva
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  Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COUNTRY: U.S.A.
                                                                                                                                                                                                                                                  US-08-461-965-307
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Indels Length 2; Xaa in location 2 is an amino acid. DB 2; . 0; Pred. No. 0; Mismatches 75.0%; Score 9; FRAGMENT TYPE: 'internal fragment 75.0 100.0%; Pre 0; Query Match 75.0 Best Local Similarity 100. Matches 1; Conservative

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Indels
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INHIBITORS AND ANTICOAGULANT PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; OTHER INFORMATION: Xaa in location 2 is an ; OTHER INFORMATION: amino acid. US-09-249-471-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        75.0%; Score 9; DB 3 100.0%; Pred. No. 0; iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                    FILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/809,455
FILING DATE: APRILI 17, 1997
APPLICATION NUMBER: PCT/US95/13231
FILING DATE: October 17, 1995
APPLICATION NUMBER: 08/486,399
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/486,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: October 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,156
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Search completed: February 5, 2001, 12:01:53 Job time: 335 sec
                                                                                                                                                                                                                                   COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 MD
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                   COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1 CURRENT APPLICATION DATA: APPLICATION NÜMBER: US/09/249,471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   REFERENCE/DOCKET NUMBER: 216/270 TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear FRAGMENT TYPE: internal fragment
                                                                                 ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       SEQUENCE CHARACTERISTICS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             2 amino acids
                                         NUMBER OF SEQUENCES: 3
                                                                                                                                               CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
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                                                                                                                                                                                                               ZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             LENGTH:
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                                                                                                                                        Sequence 52, Application US/08818253
Fatent No. 5998204
GENERAL INFORMATION:
TITLE OF INVENTION: FUORESCENT PROTEIN SENSORS FOR TITLE OF INVENTION: PETECTION OF ANALYTES
CORRESPONDENCES: 61
CORRESPONDENCES: 61
CORRESPONDENCE FISH & RICHARGON P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
STATE: CA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2; Length 2;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER: 15M COMPADIED.

OPERATING STSTEM: Windows 95
SOFTWARE: FastSEQ for Windows version 2.0b
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE: 14-MAR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
FILING DATE:
APPLICATION NUMBER: 38,347
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07257/043001
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECHONE: 619/678-5070
TELEFRAX: 619/678-5070
INFORMATION FOR SEQ ID NO: 52:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 mailto acids
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Laroche, Yves Rene
Laspers, Laurent Stephane
Gansemans, Yannick Georges Jozef
Moyle, Matthew
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100.0%; Pred. No. 0;
ive 0; Mismatches
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APPLICANT: Stanssens, Patrick Eric Hugo
APPLICANT: Messens, Joris Hilda Lieven
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 307, Application US/09249471 Patent No. 6040441 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  COMPUTER: IBM Compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 92037
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskette
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ; MOLECULE TYPE: peptide US-08-818-253-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: US
ZIP: 92037
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US-09-249-471-307
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                OM protein - protein search, using sw model
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5, 2001, 11:57:43; Search time 34.92 Seconds (without alignments) 7.778 Million cell updates/sec February

US-09-687-267-1 12

1 CXXX 4 Title: Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

195891 seqs, 67900655 residues Searched:

195891 Total number of hits satisfying chosen parameters:

seq length: 0 seq length: 2000000000 Minimum DB s Maximum DB s

Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries

Database :

PIR_66:*
1: pir1:*
2: pir2:*
3: pir3:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	R-phycoerythrin al	metallothionein-A	Iq mu chain V reqi	lidibin -	thr	R-phycoerythrin qa		-н	hemoglobin, extrac	lipopeptide WS1279	R-phycoerythrin be	protamine P1 - gor	Y protein - human	Na+/K+-exchanging	glycoprotein compo	MHC H2-K-k cell su	MHC H2-L antigen -	T-cell receptor be	T-cell receptor ga	laminin B1 - weste	hypothetical TCL3	Ig heavy chain V r	venom heptapeptide	hypothetical prote	mabinlin II chain	vicilin 57K chain	cadmium-binding he	acetylcholinestera	formylglycinamide
SUMMARIES	OI.	A22565	I51049	S43959	S55238	B22565	F22565	A33882	B45525	S65726	JU0355	C22565	137027	137263	S29881	H48394	I67345	I65546	PT0652	F41946	149421	179564	PH1408	A58512	S08606	S38516	3481	3388	340	A12016
	DB	7	~	~	~	~	N	7	~	7	~	~	7	~	~	~	~	a	~	~	~	7	~	7	7	N	~	N	N	~
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Iq H chain V-D-J r	T-cell receptor be	hypothetical L2 pr	Iq heavy chain V r	metallothionein is	urine qlycopeptide	L-serine dehydrata	leucinetRNA liga	acylase - Kluyvera	R-phycoerythrin ga	variant surface gl	variant surface gl	Iq H chain V-D-J r	T-cell receptor al	gene Cftr protein	T-cell receptor be	
PH1602	PH0932	156695	PH1407	S59622	XGHUEU	A25836	PC1002	S19288	A37521	C61512	D61512	PH1618	PH0803	157018	PH0934	
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ALIGNMENTS

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R-phycocrythrin alpha-1 chain - red alga (Gastroclonium coulteri) (fragment) C;Species: Gastroclonium coulteri C;Species: Gastroclonium coulteri C;Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993 C;Accession: A22565 R;Klotz, A.V.; Glazer, A.N. A; Klotz, A.V.; Glazer, A.N. A;Title: Characterization of the bilin attachment sites in R-phycocrythrin. A;Reference number: A22565; MuID:85182601 A;Molecule type: protein A;Residues: 1-3 KLO>

ö Gaps ó 0; Indels Length 3; DB 2; 75.0%; Score 9; DB 2 100.0%; Pred. No. 0; tive 0; Mismatches Best Local Similarity 100. Matches 1; Conservative Query Match

1 C 1 ö g

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metallothionein-A - rainbow trout (fragment)
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Species: 13-59-1996 \$sequence_revision 13-59-1996 \$text_change 21-Jul-2000
C;Accession: 151049
R;Olsson, P.E.; Kling, P.; Erkell, L.J.; Kille, P.
Eur. J. Blochem. 230, 344-349, 1995
A;Titles: Structural and functional analysis of the rainbow trout (oncorhyncus mykiss)
A;Reference number: 151049; MUD:95324545

A.Status: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA
A;Residues: 1-4 <OLS>
A;Cross-references: EMBL:X80181; NID:g1019799; PIDN:CAA56466.1; PID:g4379328

Gaps ő 0; Indels Length 4; Ouery Match 75.0%; Score 9; DB 2; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches

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RESULT S43959

Gaps

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C;Accession: A33882
C;Accession: A33882
R;Jackson, P.J.; Unkefer, C.J.; Doolen, J.A.; Watt, K.; Robinson, N.J.
Proc. Natl. Acad. Sci. U.S.A. 84, 6619-6623, 1987
A;Title: Poly(gamma-glutamylcysteinyl)glycine: its role in cadmium resistance in plan
A;Reference number: A94182; MUID:88016144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  actin I - malaria parasite (Plasmodium falciparum) (fragments)
C;Species: Plasmodium falciparum
C;Sate: 03-Jun-1993 #sequence_revision 28-Oct-1994 #text_change 09-Jun-2000
C;Accession: B45525 :
R;Wesseling, J.G.; Snijders, P.J.F.; van Someren, P.; Jansen, J.; Smits, M.A.: School. Biochem Parasitol. 35, 167-176, 1989
A;Title: Stage-specific expression and genomic organization of the actin genes 25
A;Reference number: A45525; MUID:89364996
A;Accession: B45525
      R-phycoerythrin gamma-A chain - red alga (Gastroclonium coulteri) (fragment) C; Species: Gastroclonium coulteri C; Species: Gastroclonium coulteri C; Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993 C; Accession: F22565 R; Klotz, A.V.; Glazer, A.N. J. Biol. Chem. 260, 4856-4863, 1985 A; Title: Characterization of the billin attachment sites in R-phycoerythrin. A; Reference number: A22565; MUID:85182601
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           cadmium-binding pentapeptide - downy thornapple
C;Species: Datura innoxia (downy thornapple)
C;Date: 21-May_1990 #sequence_revision 21-May-1990 #text_change 18-Jun-1993
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A,Molecule type: DNA
A;Residues: 15 < 4WES>
A;Cross-references: GB:J03988
A;Note: the authors translated the codon GAA for residue 3 as C;Comment: The actin I gene contains no introns.
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A; Molecule type: protein
A; Residues: 1-5 <JAC>
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Best Local Similarity
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C;Species: Triatcoma pallidipennis (assassin bug)
C;Species: Triatcoma pallidipennis (assassin bug)
C;Date: 19-Mar-1997 #sequence_revision 18-Jul-1997 #text_change 19-May-2000
C;Accession: S55238
R;Haendler, B.; Becker, A.; Noeske-Jungblut, C.; Kraetzschmar, J.; Donner, P.; Schleunin Bicchem. J. 307, 465-407, 1995
A;Title: Expression of active recombinant pallidipin, a novel platelet aggregation inhib A;Reference number: S55238; MUID:95251610
A;Accession: S55238
A;
                                                                                                                                                      κ.
Ig mu chain V region (clone 13) - human (fragment)
C; Species: Homo sapiens (man)
C; Date: 20-0cr1994 #sequence_revision 26-Jul-1996 #text_change 17-Mar-1999
C; Accession: S4359
R; Wagner, S.D.; Williams, G.T.; Larson, T.; Neuberger, M.S.; Kitamura, D.; Rajewsky, Nucleic Acids Res. 22, 1389-1393, 1994
Nucleic Acids Res. 22, 1389-1393, 1994
A; Reference number: S43956; MuID:94248036
A; Reference number: S43956; MuID:94248036
A; Molecule type: DNA
A; Molecule type: DNA
A; Residues: 1-4 < WRG>
C; Keywords: immunoglobulin
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B22565
R-Phycocrythrin alpha-2 chain - red alga (Gastroclonium coulteri) (fragment)
R-Phycocrythrin alpha-2 chain - red alga (Gastroclonium coulteri)
C:Species: Gastroclonium coulteri
C:Species: Gastroclonium coulteri
C:Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
C:Accession: B22565
R:Klotz, A.V.; Glasconered as Section 07-Mar-1988 #text_change 23-Mar-1993
T: Biol. Chem. 260, 4856-4865, 1985
A:Title: Characterization of the bilin attachment sites in R-phycocrythrin. A:Reference number: A22565; MUID:85182601
A:Accession: B22565
A:Molecule type: protein
A:Residues: 1-5 < KLO>
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100.0%; Pred. No. 0;
tive 0; Mismatches
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llarity 100.0%; Pred. No. 0;
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RESULT F22565

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Y protein - human (fragment)
(S.Speciess: Homo aspiens (man)
(S.Speciess: Homo aspiens (man)
(S.Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 05-Nov-1999
(S.Accession: 137263
(S.Machor, G.; Habboner, J.F.
Endocrinology 131, 2010-2015, 1992
A.Title: Novel testis germ cell-specific transcript of the CREB gene contains an alterakReference number: 137263; MUID:93010691
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S29881
C. Species: Canis lupus familiaris (dog)
C. Species: Lais lupus familiaris (dog)
C. Species: 29881
C. Accession: S29881
C. Accession: S2988
                                                                                                                                                                                                                                                       protamine PI - gorilla (fragment)
C;Species: Gorilla gorilla (gorilla)
C;Species: Gorilla gorilla (gorilla)
C;Species: Gorilla gorilla (gorilla)
C;Accession: 137027
R;Queralt, R.; Oliva, R.
Gene 133, 197-204, 1993
A;Title: Identification of conserved potential regulatory sequences of the protamine-A;Reference number: 137013; MUID:94040810
A;Accession: 137027
A;Stetus: preliminary; translated from GB/EMBL/DDBJ
A;Molecule type: DNA:
A;Residues: 1-6 <RES>
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C; Genetics:
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A; Status: preliminary; translated from GB/EMBL/DDBJ
A; Molecule type: DNA
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JU0355
Lipopeies: Streptomyces willmorei
C;Species: Streptomyces willmorei
C;Species: Streptomyces willmorei
C;Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 20-Apr-2000
C;Accession: JU0355
R;Tsuda, Y; Okada, Y; Tanaka, M.; Shigematsu, N.; Hori, Y.; Goto, T.; Hashimoto, M.
Chem. Pharm. Bull. 39, 607-611, 1991
A;Title: Structure and synthesis of an immunoactive lipopeptide, WS1279, of microbial or A;Reference number: JU0355, MUID:91300586
A;Accession: JU0355
A;Molecule type: protein
A;Residues: 1-6 <TSU>A;Note: the structure was confirmed by synthesis
C;Keywords: blocked amino end; lipoprotein
F;1/Modified site: sn-2,3-diacylglycerol (Cys) (covalent) #status experimental
F;1/Modified site: fatty acylated amino end (Cys) #status experimental
                                                                   565726
hemoglobin, extracellular, chain b - earthworm (Lumbricus terrestris) (fragment)
c.species: Lumbricus terrestris (common earthworm)
c.species: Lumbricus terrestris (common earthworm)
c.bete: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997
C.Accession: S65726
C.Accession: S65726
B.Ochim. Blophys. Acta 1292, 273-280, M.; Hosokawa, K.
Biochim. Blophys. Acta 1292, 273-280, 1996
A.Title: Characterization of the constituent polypeptides of the extracellular hemoglobil A.Reference number: S65721; MUID:96176855
A.Accession: S65726
A.Accessio
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R-phycoerythrin beta-1 chain - red alga (Gastroclonium coulteri) (fragment) C;pecies: Gastroclonium coulteri
C;pecies: Gastroclonium coulteri
C;pacession: C22565

R;Klotz, A.V.; Glazer, A.N.
J; Biol. Chem. 260, 4856-4863, 1985

A;Title: Characterization of the bilin attachment sites in R-phycoerythrin. A;Reference number: A22565; MUID:85182601

A;Molecule type: protein
A;Residues: 1-6 < KLO>
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C:Keywords: ATP; heterodimer; hydrolase; ion transport; osmoregulation; phosphoprotein;
F:4/Active site: Asp (aspartylphosphate intermediate) #starus predicted
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RESULT 15
H48394
glycoprotein component 16/major fat-globule membrane protein/MFG-E8 homolog - bovine (fr
C;Species: Bos primigenius taurus (cattle)
C;Species: Bos primigenius taurus (cattle)
C;Date: 19-Nov-1993 #sequence_revision 18-Nov-1994 #text_change 07-Feb-1997
C;Accession: H48394
R;Mather, I.H.; Banghart, L.R.; Lane, W.S.
Bjochem. Mol. Biol. Int. 29, 545-554, 1993
A;Title: The mandor fat-globule membrane proteins, bovine components 15/16 and guinea-pig
II-like sequences.
A;Reference number: A48394; MUID:93250576
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-6 <AMAT.
A;Reperimental source: milk
A;Reperimental source: milk
A;Note: sequence extracted from NCBI backbone (NCBIP:131518)
C;Keywords: glycoprotein

0; Gaps 0; Indels Query Match 75.0%; Score 9; DB 2; Length 6; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches 0; Indel.

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Search completed: February 5, 2001, 12:02:32 Job time: 289 sec

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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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OM protein - protein search, using sw model

February 5, 2001, 12:01:23; Search time 20.72 Seconds (without alignments) 6.234 Million cell updates/sec Run on:

US-09-687-267-1 12 1 CXXX 4 Title: Perfect score: Sequence:

Scoring table:

BLOSUM62 Gapop 10.0 , Gapext 0.5

88757 Total number of hits satisfying chosen parameters:

88757 segs, 32294092 residues

Searched:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_39:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	P80709 carcinus ma	P02729 homo sapien	6 carci	548	couns /		-	P29177 bos taurus						eisenia		oryc	P42994 raja clavat		O54296 salmonella	P24047 stomopneute	pichia	pich		chely		P50983 conus imper	O22426 lotus japon	369 sesban		207	371	122	PO4559 catostomus
SUMMARIES	ΩI	ACT_CARMA	GLUR_HUMAN	CCAP_CARMA	CONO_CONGE	CONO_CONST	DNF1_LOCMI	ISOT_CYPCA	MGMT_BOVIN	OXYA_SCYCA	OXYA_SQUAC	OXYF_SCYCA	OXYT_BUFRE	OXYT_CYPCA	OXYT_EISFO	OXYI_OCTVU		- 1	OXYV_SQUAC	RS11_SALTY	SAP_STOVA	TAL1_PICJA	TAL3_PICJA	GLEM_HUMAN	GON2_CHEPR	RCA_PINPS	CXA1_CONIM	NO40_LOTJA	- 1	- 1	RR16_GINBI	- 1	ı	UR2B_CATCO
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	Result No.	П	7	m	4	S	9	7	ω	6	10	11	12	13	14	15	16		18	19	20	21	22	23	24	25	26	27	28	29	30		32	33

P04561 cyprinus ca P01147 gillichthys P811022 polyodon sp P35490 scyliorhinu P17776 escherichia P15987 glycine max P15987 glycine max P15987 glycine max P15987 conus stria P28878 conus stria P28878 conus texti P28878 hordeum vul P28525 hordeum vul			ı crab). Malacostraca; emata; Brachyura; inus.	E., Webster S., steroidogenesis.";	INS THAT ARE IN BIQUITOUSLY EXP	D PI OF THIS PROTEIN IS:	CRC64;	Length 8; 0; Indels 0; Gaps C	rata; Euteleostomi; nidae; Homo.
1 UR2B_CYPCA 1 UR2_GILMI 1 UR2_BOLSP 1 UR2_SCYCA 1 Y2PY_ECOLI 1 ACT7_SCYBN 1 CXA1_CONST 1 CXA2_CONGE 1 CXA2_CONGE 1 CXA2_CONGE 1 CXA2_CONGE 1 CXA2_CONGE 1 CXA2_CONGE 1 GER1_HORVU 1 GER2_HORVU	ALIGNMENTS	ESULT 1 CI_CARMA D ACT_CARMA STANDARD; PRT; 8 AA. C P80709; I 01-0CT-1996 (Rel. 34, Created) I 01-0CT-1999 (Rel. 34, Last sequence update) I 15-JUL-1999 (Rel. 38, Last annotation update)	ihore crab) (Green cral copoda; Crustacea; Mala Decapoda; Pleocyemata Portunidae; Carcinus	rpentier G., Granjeon me implicated in crab	HIGHLY CONSERVED PROTEINS SELL MOTILITY AND ARE UBIOU	LOSTOPLASMIC. E 2D-GEL THE DETERMINED DA	DR PROSITE; PS00405; ACTINS_1; PARTIAL. DR PROSITE; PS00432; ACTINS_2; PARTIAL. DR PROSITE; PS01132; ACTINS_2; PARTIAL. KW Structural protein. FT NOW_TER 8 8 SC SEQUENCE 8 AA: 976 MW; 1424005AB2CAAEB3 CRC	%; Score 9; DB 1; 1 .0%; Pred. No. 0; 0; Mismatches	JARD; PRT; 8 AA. 1, Created) 1, Last sequence update) 2) Last annotation update) 3) Chordata; Craniata; Vertebrata; Primates; Catarrhini; Hominidae;
3 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2 2		STANDARD; kel. 34, Cre kel. 34, Las kel. 38, Las	TT). ts (Common stazoa; Arthu t; Eucarida;	Somme G., Ca D.; Nse. An enzy	ACTINS ARE TYPES OF	AALOCATIONS SOUS: ON THE W IS: 46 KI T: BELONGS I	105; ACTINS 132; ACTINS 132; ACTINS 5tein. 1 1 1 8 8 8 AA; 976 MW;	75.0%; larity 100.0% Conservative	DAR 11, 0, C.
22222222222		1 CT_CARWA 80709; 1-0CT-1996 (F 1-0CT-1999 (F	orin (FRAGMEN arcinus maene akaryota; Met amalacostrace abrachyura; F	1) EQUENCE. achaise F., E aghdassarian A transaldole	docrine 5:2: !- FUNCTION: IN VARIOUS	- SUBCELLANS - MISCELLANS 6.8, ITS N - SIMILARITY	WOSITE; PSOU ROSITE; PSOU ROSITE; PSOU1 ELUCTURAL PIC DN_TER SQUENCE 8 P	y Match Local Similarity hes 1, Conser 1 C 1 2 C 2	UMAN -1986 (-1986 (-2000 (GLYCOPE GLYCOP
W W W W W W A 4 4 4 4 4 4 4 4 4 4 4 4 4		RESULT ACT_CARMA ID ACT_ AC P807 DT 01-0 DT 01-0 DT 15-J	8 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0 0	RN RP SI	75008 E	300002	S T T T W S S S S S S S S S S S S S S S	Ouery M Best Loc Matches Oy 1	RESULT 2 GLUR_HUMAN ID GLUR_C AC P02725 DT 21-JUI DT 21-JUI DT 01-CTI DT 01-CTI DT 01-CTI C C EUKALY OC MAMMAI

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Cruz L.J., de Santos V., Zafaralla G.C., Ramilo C.A., Zeikus R.D.,
Gray W.R., Ollvera B.M.;
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"Invertebrate vasopressin/oxytocin homologs. Characterization of
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              "Invertebrate vasopressin/oxytocin homologs. Characterization of peptides from Conus geographus and Conus straitus venoms.", J. Biol. Chem. 262:15821-15824(1987).
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"Peptide toxins from venomous Conus snails.";
Annu. Rev. Blochem. 57:665-700(1988).
-1. SIMILARITY: BLONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; A28495, A28495.
INTERPRO; IPRO00981; -.
PROSITE; PS00220; hormone4; 1.
Hormone; Amidation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conus geographus (Geography cone).
Eukaryota; Metazoa; Mollusca; Gastropoda; Caenogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
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Conus stitatus (Striated cone).
Eukaryota; Macezoa; Mollusca; Gastropoda; Caenogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
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D4FC276EB4540059 CRC64;
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01-NOV-1988 (Rel. 09, Last sequence update)
10-NOV-1995 (Rel. 32, Last annotation update)
LTS-CONOPRESSIN:G.
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(Rel. 09, Last sequence update)
(Rel. 32, Last annotation update)
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llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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100.0%; Pre
0;
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9 AA; 1037 MW;
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Best Local Similarity
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Best Local Similarity
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01-NOV-1995 (
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P05486;
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P05487;
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SEQUENCE
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SEQUENCE.
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CONO_CONGE
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                                                  Lote C.J., Weiss J.B.;

"Identification in urine of a low-molecular-weight highly polar
glycopeptide containing cysteinyl-galactose.";
glycopeptide containing cysteinyl-galactose.";
electron. J. 123:25p-25p(1971).
electron. J. 123:25p-25p(1971).
electron. THE IDENTITY OF THE GLYCOPROTEIN FROM WHICH THIS PEPTIDE
IS DERIVED IS UNKNOWN. NO PHYSIOLOGICAL FUNCTION HAS BEEN
ATTRIBUTED. AN ERYTHROCYTE MEMBRANE GLYCOPEPTIDE HAVING A
SIMILAR STRUCTURE HAS ALSO BEEN FOUND.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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Proc. Natl. Acad. Sci. U.S.A. 84:575-579(1987).
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MEDLINE-93050243; PubMed-1426284;
Cheung C.C., Loi P.K., Sylwester A.W., Lee T.D., Tublitz N.J.;
Cheung Y.C.C., a cardioactive neuropeptide from the tobacco hawkmoth, Manduca sexta.;
FEBS Lett. 313:165-168(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         molitor and Spodoptera eridania";
Biol. Chem. Hoppe-Seyler 374:1065-1074(1993).
-!- FUNCTION: THE EFFECT OF CCAP IS BOTH INO- AND CHRONOTROPIC.
-!- TISSUE SPECIFITY: STORED IN PERICARDIAL ORGANS AND RELEASED PIR; A26363; A26363.
PIR; S27233; S27233.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Carcinus maenas (Common Shore crab) (Green crab),
Manduca sexta (Tobacco hawkmoth) (Tobacco hornworm),
Tenebrio molitor (Yellow mealworm), and
Spodoptera eridania (Southern armyworm).
Eukaryota; Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Bucarida; Decapoda; Pleocyemata; Brachyura;
Eubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Indels
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 8;
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8 AA; 855 MW; C2D87AA1F5B1EB1E CRC64;
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C5A861A9CDD44EB9 CRC64;
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MEDLINE-94176032; PubMed-8129851;
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01-OCT-1994 (Rel. 30, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
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llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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SEQUENCE.
MEDLINE-72062338; PubMed-5126885;
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DISULFID 3 9
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Best Local Similarity
Matches 1; Conserv
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CARBOHYD
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P38556;
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Pred. No. 0;
Mismatches
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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                                                                                                                     Cyprinus carpio (Common carp).
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                              STANDARD;
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DISULFID 1
MOD_RES 9
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                          ISOT_CYPCA
P42993;
                                                          01-NOV-1995
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01-NOV-1995
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P29177;
                                                                                                       ISOTOCIN.
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               SOT_CYPCA
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MGMT_BOVIN
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MEDLINE-88077077; PubMed=3689410;
Proux J.P., Miller C.A., Li J.P., Carney R.L., Girardie A.,
Delaage M., Schooley D.A.;
"Identification of an arginine vasopressin-like diuretic hormone from
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Eukaryota: Metazoa; Arthropoda; Tracheata: Hexapoda; Insecta;
Pterygota: Neoptera: Orthopteroidea; Orthoptera; Caelifera;
Acridomorpha; Acridoidea; Acrididae; Locusta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Blochem. Blophys. Res. Commun. 149:180-186(1987).
-!- FUNCTION: DIURETIC HORMONE.
-!- SUBUNIT: F2 IS AN ANTIPARALLEL DISULFIDE LINKED DIMER OF
-!- SIBULITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; A29477; A29477
peptides from Conus geographus and Conus straitus venoms."; J. Biol. Chem. 262:15821-15824(1987).
                                                                        Gray W.R., Olivera B.M., Cruz L.J.;
Peptide toxins from venomous conus snails.";
Annu. Rev. Biochem. 57:665-700(1988).
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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INTERCHAIN (WITH C-1')
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17EB176EB4540050 CRC64;
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56EB176EB451A057 CRC64;
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01-AUG-1990 (Rel. 15, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
LOCUPRESSIN (DIURFIIC NEUROPEPTIDE F1/F2).
                                                                                                                                                                                                                                                                                   DB 1;
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100.0%; Pred. No. 0;
:ive 0; Mismatches
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Mismatches
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                                                                                                                                               INTERPRO, IPRO00981; -.
PFAM; PF00220; hormone4; 1.
PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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                                                          MEDLINE=89024586; PubMed=3052286;
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Conservative 0;
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9 9
9 AA; 1031 MW;
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Best Local Similarity
Matches 1; Conserv
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Matches 1; Conserv
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01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
01-DEC-1996 (Rel. 34, Last annotation update)
METHYLATED-DNA--PROTEIN-CYSTEINE METHYLTRANSFERASE (EC 2.1.1.63) (6-0-
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INTERPRO; IPRO01497; -.
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Actinopterygii; Neopterygii; Teleostei; Buteleostei; Ostariophysi;
Cypriniformes; Cyprinidae; Cyprininae; Cyprinus.
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Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
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9 AA; 969 MW; 17FF476EB455B04B CRC64;
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                    (Rel. 32, Created)
(Rel. 32, Last sequence update)
(Rel. 32, Last annotation update)
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9 AA.
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Gaps
"Phylogeny of the neurohypophysial hormones. Two new active peptides isolated from a cartilaginous fish, Squalus acanthias.";
Eur. J. Biochem. 29:12-19(1972).
                                                                                                                     Acher R., Chauvet J., Chauvet M.-T., Fontaine M.;
"Identification of 2 new neurohypophyseal hormones, valitocin (Val8-oxytocin) and aspartocin (Asn4-oxytocin) in a selachian fish, the spiny dog-fish (Squalus acanthias).";
C. R. Acad. Sci., D. Sci. Nat. 274:313-316(1972).
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Chondrichthyes,
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyllorhinidae; Scyllorhinus.
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9 AA; 996 MW; 17F8376EB444404B CRC64;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
Hormone; Amidation.
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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100.0%; Pre
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ID OXYF_SCYCA
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Squalus acanthias (Spiny dogfish).
Elwaryota: Metazoa: Chordata: Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Squalea: Squaloidei; Squalidae; Squalus.
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Eukaryota, Metazoa, Chordata; Craniata; Vertebrata; Chondrichthyes;
Elssmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyliorhinidae; Scyliorhinus.
                                                                ALKYL GROUP ACCEPTOR (BY SIMILARITY).
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Chauvet J., Roulle Y., Chauveau C., Chauvet M.-T., Acher R.;
Special evolution of neurohypophysial hormones in cartilaginous
fishes: asvatocin and phasvatocin, two oxytocin-like peptides
isolated from the spotted dogfish (Scyliorhinus caniculus).";
Proc. Natl. Acad. Sci. U.S.A. 91:11266-11270(1994).
-i- FUNCTION: DISPLAYS OXYTOCIA ACTIVITY ON RAT UTERUS.
-i- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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17EDD76EB444404B CRC64;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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01-NOV-1995 (Rel. 32, Created)
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01-NOV-1995 (Rel. 32, Last annotation update)
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                       Transferase; Methyltransferase.
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100.0%; Pred. No. 0;
ive 0; Mismatches
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MEDLINE-73031727; PubMed-5083097;
Acher R., Chauvet J., Chauvet M.-T.;
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    PS00374; MGMT; PARTIAL
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                       DNA repair;
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P42996;
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TISSUE-NERVE ENDINGS;
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Hormone; Amidation.
DISULFID 1:
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01-NOV-1995
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P80027;
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P42998;
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OXYT_OCTVU

OXYT_OCTVU

DC P80027/D
DT 01-MAR.
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Acher R., Chauvet J., Chauvet M.-T., Crepy D.;
"Characterization of neurohypophyseal hormones from a fresh water bony fish, the carp (Cyprinus carpio). Comparison with hormones from sea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cyprinus carpio (Common carp), and Petromyzon marinus (Sea lamprey). Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Actinopterygii; Neopterygii; Teleostei; Euteleostei; Ostariophysi; Cypriniformes; Cyprinidae; Cypriniae; Cypriniae;
                                                                                                                                                                                                                                                                                                                                                                                                                               MEDITARIA SURVAINTENTED LODG.

Chauvet J., Michel G., Ouedraogo Y., Chou J., Chait B.T., Acher R.;

A new neurohypophysial peptide, seritocin ([Ser5,Ile8]-oxytocin),
identified in a dryness-resistant African toad, Bufo regularis.";
Int. J. Pept. Protein Res. 45:482-487(1995).
-!- FUNCTION: DEVOID OF OXYTOCIC ACTIVITY.
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PRAM: PF00220; hormone4; 1.
PRAM: PF00220; hormone4; 1.
PRAM: PAUSE SIMILARITY: PAUGO SITE: PS00264; NEUROHYPOPHYS_HORM; FALSE_NEG.

DISULPID
                                                                                                                                                                                                     Bufo regularis (African toad).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Amphibia; Batrachia; Anura; Neobatrachia; Bufonoidea; Bufonidae;
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MEDLINE-88225976; PubMed-3371648;
Lane T.F., Sower S.A., Rawauchi H.;
Lane T.F., Sower Dan, Rawauchi H.;
Marghine vasotocin from the pituitary gland of the lamprey (Petromyzon marinus): isolation and amino acid sequence.";
Gen. Comp. Endocrinol. 70:152-157(1988).
-!- FUKCTION: ANIIDIURETIC HORMONE.
-!- SIMILARIY: BELGONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; BG1364; BG1364.
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9 AA; 983 MW; 17FF476EA5A6D04B CRC64;
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01-NOV-1991 (Rel. 20, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
   9 AA.
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Comp. Biochem. Physiol. 14:245-254(1965).
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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   STANDARD;
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Best Local Similarity
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OXYT_CYPCA
AC P23879;
DT 01-NOV-1991 (F)
DT 01-NOV-1991 (F)
DT 01-NOV-1991 (F)
DE VASOTOCIN.
CC CURLARYOTES
CC CYPTINUS CATP
CC CYPTINUS CATP
CC CYPTINISCARP
CC CYPTINIFORMES;
RN (1]
RN SECUENCE.
RC SPECIES=C.CARI
RA ACHER R., CHARI
RA ACHER R., CHARI
RY WATER DONY FILL
RL COMP. BIOCHEM
RT FISH, THE SECUENCE.
RC SPECIES=P.MAR
RT AND SECUENCE.
RC SPECIES=P.MAR
RT RATGININE WAS
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CC -!- FUNCTION:
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   OXYT_BUFRE P42995;
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Biochem. Biophys. Res. Commun. 198:393-399(1994).
-! FUNCTION: POTENTIALES SPONTANEOUS CONTRACTIONS OF THE GUT AND ALSO PULSATORY CONTRACTIONS AND BLADDER-SHAKING MOVEMENT OF THE NEPHRIDIA. MAY BE INVOLVED IN OSMOREGULATION OF THE ANIMAL THROUGH NEPHRIDIAL. EUNCTION.
-! SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; PC2021; PC2021.
                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eisenia foetida (Common brandling worm) (Common dung-worm).
Eukaryota; Metazoa; Annelida; Clitellata; Oligochaeta; Haplotaxida;
Lumbricina; Lumbricidae; Eisenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   MEDLINE-94121660; Pubmed-8292046;
Oumi T., Ukena K., Matsushima O., Ikeda T., Fujita T., Minakata H.,
Nomoto K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               "Annetocin: an oxytocin-related peptide isolated from the earthworm,
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Incirrata; Octopodidae; Octopus.
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9 AMIDATION.
1053 MW; 17EB176EB456D04B CRC64;
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D4EEB76EB45412C9 CRC64;
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(Rel. 32, Last sequence update)
(Rel. 32, Last annotation update)
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01-WAR-1992 (Rel. 21, Last sequence update)
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100.0%; Pred. No. v.
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100.0%; Pred. No. 0;
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                                                                                                Thereof is the control of the oxytocin/vasopressin family isolated from nerves of the cephalopod Octopus vulgaris.";

In nerves of the cephalopod Octopus vulgaris.

In nerves of the cephalopod Octopus vulgaris.

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MEDLINE=92270139; PubMed=1589145; Reich G.;
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Search completed: February 5, 2001, 12:06:02 Job time: 279 sec

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Q1599 homo sapten
Q94887 homo sapten
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Q69473 human herpe
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O12102 caprine art
O12103 experine art
O12104 caprine art
O12107 arcobacteri
Q47475 escherichia
Q94865 klebsiella
Q94845 escherichia
Q94813 escherichia
Q913118 homo sapien

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SECUENCE FROM N.A. MEDLINE=93010691; PubMed=1396344; MEDLINE=93010691; PubMed=1396344; Meaber G., Habener J.F.; Movel testis germ cell specific transcript of the CREB gene contains an alternatively spliced exon with multiple in-frame stop codons."; Endocrinology 131:2010-2015(1992). EMBL: X68994; CAA48700.1; NON_TER
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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Homo sapiens (Human).
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066113 cherry leaf
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02564 fugu rubrip
032560 escherichia
01588 homo sapien
015890 homo sapien
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Viruses; ssRNA positive-strand viruses, no DNA stage; Comoviridae;
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Thesis (1992), Biologia Molecular y Virologia Vegetal, CIT-INIA.
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"Long, nearly identical untranslated sequences at the 3' terr
regions of the genomic RNAs of cherry leafroll virus (walnut
strain).";
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EMBL; Z34265; CAA84019.1; -.
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Rehm B.H.A., Ertesvag H., Valla S.;
"A new Azotobacter vinelandii mannuronan C-5-epimerase gene (algG) is part of an alg gene cluster physically organized in a manner similar to that in Pseudomonas aeruginosa.";
J. Bacteriol. 178:5884-5889(1996).
EMBL; X87973; CAA61230.1; -.
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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                           Azotobacter vinelandii.
Bacteria; Proteobacteria; gamma subdivision; Pseudomonadaceae;
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01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
0RPHAN RECEPTOR TR4-NS (FRAGMENT).
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MEDLINE=96198747; PubMed=8612486;
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MEDILINE-81001992; bubMed=7407922;

Dhar R., Chanock R.M., Lai C.J.;

"Nonviral oligonucleotides at the 5' terminus of cytoplasmic influenza viral mrNA deduced from cloned complete genomic sequences.";

Call 21:495-500(1980)

EMBL; M25045; AAA43202.1; -. Gaps ö Q67113; 01-NOV-1996 (TrEMBLrel. 01, Created) 01-NOV-1996 (TrEMBLrel. 01, Last sequence update) 01-NOV-1998 (TrEMBLrel. 08, Last annotation update) (UDORN/72) HEMBAGGUTININ (SEG 4) CDNA, 3' END (FRAGMENT). Influenza A virus. Viruses; ssRNA negative-strand viruses; Orthomyxoviridae; Influenza virus A and B group. 75.0%; Score 9; DB 12; Length 7; llarity 100.0%; Pred. No. 0; Conservative 0; Mismatches 0; Indels 834 MW; 605EB0544EA40030 CRC64; m · U

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Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
NCBI_TaxID=85755;
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STRAIN=GOMEN,
Crawford-Miksza L.K., Nang R.N., Schnurr D.P.;
"Molecular surveillance of strain variation in adenoviruses acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065065; AAD03662.1; -.
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STRAIM=KN T96-0620, S-1058, CL 68578;
Crawford-Miksza L.K., Nang Kn., Schnurr D.P.;
Crawford-Miksza L.K., Nang R.N., Schnurr D.P.;
Wolecular surveillance of strain variation in adenoviruses acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL, AF065069; AAD03668.1;
EMBL, AF065066; AAD03664.1;
EMBL, AF065067; AAD03666.1;
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MEDLINE=97442476; PubMed=9295353;
Plummer N.W., McBurney M.W., Melsler M.H.;
Plummer N.W., McBurney M.W., Melsler M.H.;
Alternative splicing of the sodium channel SCN8A predicts a truncated two-domain protein in fetal brain and non-neuronal cells.";
J. Baol. Chem. 272:24008-24015(1997).
EMBL: U97673; AAB80916.1; --
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Crawford-Miksza L.K., Nang R.N., Schnurr D.P.;
"Molecular surveillance of strain variation in adenoviruses causing acute respiratory disease, AV 4 and AV 7a.";
submitted (MAX-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065064; AAD03659.1; -.
EMBL; AF065062; AAD03659.1; -.
                                                                                                                              Human adenovirus type 4.
Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
NCBI_TaxID=28280;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fugu rubripes (Japanese pufferfish) (Takifugu rubripes).
Bukaryota: Metazoa; Chordata; Cranhata: Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        01-JAN-1998 (TrEMBLrel. 05, Created)
01-JAN-1998 (TREMBLrel. 05, Last sequence update)
01-MOV-1998 (TrEMBL-el. 08, Last annotation update)
TRUNCATED VOLFAGE-GATED SODIUM CHANNEL ALPHA SUBUNIT (FRAGMENT).
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 7;
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01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
PVI CORE PROTEIN (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                NON_TER 1 1 1 SEQUENCE 7 AA; 730 MW; 75B72EA2C73772A0 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                        NON_TER 1 1 SEQUENCE 7 AA; 980 MW; 7B5EA414140322A0 CRC64;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                12;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       7 AA
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Pred. No. 0;
Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Score 9;
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Created)
                                                                                                                                                                                                                              SEQUENCE FROM N.A. STATE 67, 55142; STRAIN-Z-G 95-873, RI-67, 55142;
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100.0%; Pre
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Gaps

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0; Indels

Length 8;

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TISSUB-PLACENTA;
Lee C.C., Yazdani A., Wehnert M., Bailey J., Couch L., Xiong M.,
Coolbaugh M.I., Chinault C.A., Baldini A., Lindsay B.A., Zhao Z.Y.,
                   Homo sapiens (Human).
Eukaryota; Metasoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                        8 AA; 975 MW; 605EA6C5BEA5A2D3 CRC64;
                                                                                                                                                                                                                                                                                                                                              75.0%; Score 9; DB 4
100.0%; Pred. No. 0;
ative 0; Mismatches
 (CLONE XP19G12A) (FRAGMENT).
                                                                                                                                                                                       Caskey C.T.H.;
Hum. Mol. Genet. 0:0-0(0).
EMBL; L32083; AAA73880.1;
NON_TER
                                                                                                                                                                                                                                                                                                                                                                                      1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               PRELIMINARY;
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Best Local Similarity
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Best Local Similarity
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TISSUE=PLACENTA;
                                                                                                                 SEQUENCE FROM N.A.
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                                                                         NCBI_TaxID=9606;
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NON_TER
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Q15900
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Homo sapiens (Human).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
                                                                         Escherichia coll.
Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
Escherichia.
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                                                                                                                                                                                                                                                   [2]
SEQUENCE FROM N.A.
SEQUENCE FROM N.A.
HESSLINGER C., Fairhurst S.A., Sawers G.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ001620; CAA04875.1; -.
                                                                                                                                                                                                           Hesslinger C., Sawers G.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases
                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 8;
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Last sequence update)
Last annotation update)
01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-AUG-1998 (TrEMBLrel. 07, Last annotation update)
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01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-AUG-1998 (TrEMBLrel. 07, Last annotation update)
(CLONE XP15H8A) (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                8 AA; 1068 MW; 0315A37EAB5B0763 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE 8 AA; 1000 MW; 3A505EB044140DC4 CRC64;
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0. 0;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           8 AA.
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100.0%; Pred. No. 0;
Live 0; Mismatches
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                                     PROPIONATE KINASE (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Hum. Mol. Genet. 0:0-0(0).
EMBL; L32069; AAA73878.1; -.
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Best Local Similarity
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                                                                                                                                 NCBI_TaxID=562;
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                                                                                                                                                                                             STRAIN=W3110;
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01-NOV-1996 (
01-AUG-1998 (
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SEQUENCE
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Q15890
ID Q15890
AC Q158900
DT 01-NOV
DT 01-AUG
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Matches
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Q15888
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                                                                                                                                                                                Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 8;
                                            01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NUV-1996 (TrEMBLrel. 07, Last annotation update)
(CLOME XYPHIA) (FRAGMENT).
HOMO Saplens (Human).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         01-NOV-1999 (TrEMBLrel. 12, Created)
01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
NUCLEAR LIM INTERACTOR (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  8 AA; 931 MW; B5DDC403369AAEB1 CRC64;
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8 A.A.
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100.0%; Pred. No. 0;
tive 0; Mismatches
PRT;
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"Evidence for insufficient chondrocytic differentiation during repair of full-thickness defects of articular cartilage.";
Matrix Biol. 15:39-47(1996).
BMBL: S83371; AAD14433.1; -.
NON_TER
SEQUENCE 8 AA; 1028 MW; B859C7272EA77371 CRC64;
                                                                   SEQUENCE FROM N.A.
Drechsler M., Schumacher V., Friedrich S., Wildhardt G., Giesler S., Schroth A., Bodem J., Royer-Pokora B.;
"Genomic structure, alternative transcripts and chromosomal localization of the human LIM domain binding protein I gene
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Gaps
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-MAY-1999 (TrEMBLrel. 04, Last sequence update)
PRO ALPHA 1 TYPE III COLLAGEN PROTEIN (FRAGMENT).
Oryctolagus cunfculus (Rabbit).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Butheria; Lagomorpha; Leporidae; Oryctolagus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       [1]
SEQUENCE FROM N.A.
MEDLINE=96377339; PubMed=8783186;
Metsaranta M., Kujala U.M., Pelliniemi L., Osterman H., Aho H.,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                            LDB1/NLI.";
Submitted (JUN-1999) to the EMBL/GenBank/DDBJ databases.
EMBL, 121243097; CAB45408.1;
NON_TER 8
SEQUENCE 8 AA; 767 MW; EEGEBDDEB862D5B6 CRC64;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match 75.0%; Score 9; DB 4; Length 8; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches 0; Indel.
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100.0%; Pred. No. 0;
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        PRT;
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Best Local Similarity
Matches 1; Conserva<sup>†</sup>
NCBI_TaxID=9606;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5
C 5
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002831
D 002831
D 002831,
DT 01-JUL.
DT 01-MAY.
DT 01-MAY.
DC C MAMMAI.
OC MAM
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Search completed: February 5, 2001, 12:05:38 Job time: 289 sec

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Gaps

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0; Indels

Conservative

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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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OM protein - protein search, using sw model

Run on:

5, 2001, 12:01:19; Search time 42.35 Seconds (without alignments) 3.230 Million cell updates/sec February

US-09-687-267-4

1 XCXX 4 Title: Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

268485 seqs, 34193795 residues Searched:

Total number of hits satisfying chosen parameters:

268485

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

Database :

| SIDSI/gcgdata/geneseq/geneseqp/AA1980.DAT:*
| SIDSI/gcgdata/geneseq/geneseqp/AA1981.DAT:*
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/SIDS1/gcgdata/geneseq/geneseqr/AA1989.DAT: /SIDS1/gcgdata/geneseq/geneseqp/AA1990.DAT: /SIDS1/gcgdata/geneseq/geneseqp/AA1991.DAT: /SIDS1/gcgdata/geneseq/geneseqp/AA1991.DAT:

/SIDS1/gcgdata/geneseq/geneseqp/AA1997.DAT:*/SIDS1/gcgdata/geneseq/geneseqp/AA1998.DAT:*/SIDS1/gcgdata/geneseq/geneseqp/AA1999.DAT:*/SIDS1/gcgdata/geneseq/geneseqp/AA1999.DAT:*/SIDS1/gcgdata/geneseq/geneseqp/AA2000.DAT:* /SIDS1/gcgdata/geneseq/geneseqp/AA1993.DAT: /SIDS1/gcgdata/geneseq/geneseqp/AA1994.DAT: /SIDS1/gcgdata/geneseq/geneseqp/AA1995.DAT: /SIDS1/gcgdata/geneseq/geneseqp/AA1996.DAT:

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

SUMMARIES

		Description	Sequence which cor	Binding receptor w	Synthetic peptide	Metal trapping pep	Metal trapping pep	IGF-1 analoque N-t	IGF-1 analogue N-t	IGF-1 analogue N-t	Human interleukin-	Anti-inflammatory	Anti-inflammatory	Anti-inflammatory
		dī	P10301	P90982	P91660	R34022	R34023	R51439	R51440	R51441	W19847	W56231	W56211	W56200
		h DB	3	3 10	3 10	3 14	3 14	3 15	3 15	3 15	3 18	3 19	3 19	3 19
		Lengt												
æ	Query	Match Length DB	75.0	75.0	75.0	75.0	75.0	75.0	75.0	75.0	75.0	75.0	75.0	75.0
		Score	0	O	σ	σ	σ	σ	6	σ	σ	o.	6	თ
	Result	No.	Ħ	7	m	4	S	Ø	7	ω	σ	10	11	12

flamma d	used in	rrz3	e ot	e of	lin	ic a		e of	oŧ	of	enc	-	Peptide fragment #	Peptide fragment #	loxin	plate	platelet a	bressure r	7F.9	Farnesyl-protein :	÷	å	÷	yl-prote	y1-	÷	arnesyl-prote	Farnesyl-protein t	•	å	•	Fluorophore-labell
W56172	WGG057	X51896	10138	P10093	P10382	P20217	P30083	P30085	P40538	P40539	70833	P91661	R11077	R11079	R11508	R11744	74	5	R13935	R14723	R15751	ī	-	'n	R15755	R15756		R15758	R15759	576	Ö	Y80840
13		77				3	4			2					12	12	12	12	12	12	12	12	12	12	12	12	12	12	12	12	12	21
m (n (η.	4	4	7	4	4	4	4	7	4	4	7	4	4	4	4	4	4	7	4	7	7	4	4	4	4	4	4	7	4	4
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75	- 1	7 1		7.5	75	75	7.5	75	75	7	7.5	7.5	7.5	75	7.5	75	75	7.5	75	75	7.5	75	7.5	7	7.5	7.5	7.5	7.5	75	7.5	75	75
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13	# L	T ,	16	17	18	19	20	21	22	23	24	22	56	27	28	53	30	31	32	33	34	35	36	37	38		40	41	42	43	44	45

ALIGNMENTS

P10301 ID P10301 standard; Protein; 3 AA. P10301; RESULT

(first entry) 19-AUG-1992

Sequence which corresp. to resides 19-21 of insulin A chain.

Cystine peptide; insulin activity; hormone; diabetes therapy; antidiabetic agent.

Mammal.

Location/Qualifiers Key Disulfide-bond

/note= "bonded to Cys(5) of P10115, which is residue 19 of the B chain fragment B15-27"

DD147942-A.

29-APR-1981

30DD-0011868 01-FEB-1980;

79DD-0215100 20-AUG-1979;

(LOSS/) LOSSE G.

Stange H; Losse G,

WPI; 1981-53498D/30 (53498D).

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Gaps

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Indels

us-09-687-267-4.rag

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The inventors claim synthetic peptides which correspond to sequences occurring in EGF, but excluding EGF. The peptides are angiogenic and have corresp. applications, eg for the healing of wounds and burns. Their relative shortness means that they pose fewer synthesis problems than the entire EGF molecule. They can be admin. singly or association with each other or in association with an angiogenic
                                                                                                                                                                                                                                                                  to residues 12-14 of naturally occurring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Synthetic peptide active in stimulating angiogenesis -
has sequences corresponding to amino acid sequences occuring in
                                                                                                                                                                                                                                                                                                                     Epidermal growth factor; angiogenesis; synthetic peptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                             /label-OTHER
/note="(aceto amido methyl) NH2-Cys"
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100.0%; Pre
0;
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ö
                                                                                                                                                           P91660 standard; protein; 3 AA.
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                                                                                                                                                                                                                                                                Synthetic peptide corresp. tc epidermal growth factor (EGF)
                                                                                                                                                                                                                                                                                                                                                                                         /label=OTHER
/note="H-Gly
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       88WO-AU00300
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          87AU-0003629
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 Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      has sequences correspondepidermal growth factor.
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ID R3
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    The article for inactivating a toxic material comprises a solid carrier bearing a first receptor which binds the target ligand and a second receptor which degrades the target ligand. This synthetic peptide is a preferred first receptor. The article may be used for covering surfaces to protect or decontaminate the surface. The article is especially toxic organic cyds., esp. organophosphorous cyds., (eg pesticides and nerve poisons, bacteria and viruses, in environmental, chemical, military and industrial settings.
                                                                                      The inventors claim the prepn. of new cysteine peptides with insulin-
                                                                                                      like activity. The new peptides comprise a sequence with corresp. to a central portion of the insulin B chain (esp. B15-27) and a cysteine-contg. peptide with an AA sequence corresp. to a terminal portion of the insulin A chain (esp. A1-7). The new cysteine peptides are antidiabetic agents with appreciable insulin activity (lower than that of insulin itself) and low antigenic activity.
                                                                                                                                                                                                                                                                                                                       Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ..... material; inactivator; organophosphorous cpds.; nerve poison; pesticide; decontaminant; military.
Cysteine peptide(s) with antidiabetic activity - contg. fragments of insulin A and B chains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Binding receptor with selectivity for a target ligand, borne by an article for inactivating toxic materials.
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100.0%; Pred. No. 0;
iive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                               P90982 standard; peptide; 3 AA.
                                                   Claim 1; Page 10; 11pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    88WO-US03422.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                 New acyclic peptide with metal trapping activity - exhibits in vivo metal transporting action and can be used as drug, diagnostic agent or functional material
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The acyclic peptide (see R31340) with metal trapping activity produced by fragment condensation using the peptide fragments R34022-29.
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in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
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100.0%; Pred. No. 0;
Live 0; Mismatches
                                              ./note= "OMe protected"
of cysteine"
                                                                                                                                                                                                                                                                                       (ICHI-) ICHIKAWA GOSEI KAGAKU KK.
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Walton PE;
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(GROP-) GROPEP PTY LTD.
(USDA ) US SEC OF AGRIC.
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 AA;
                        Modified-site
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                                                                                                                                                                                          24-MAY-1991;
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R51439
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                                                                                                                                                                                                                                                                                                                                   /note= "Cys(Acm); Acm= acetamidomethyl-protected SH
   of cysteine"
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                                                                                                                   Electrostatic interaction; metalloprotein; binding site;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
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llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                    /note= "Boc protected"
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/note= "OMe protected"
                                                                    Metal trapping peptide intermediate.
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Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
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                                                            Growth of birds is increased by (a) admin. to the bird, in ovo, insulin-like growth factor (IGF)-1 or -2 or their active analogues; (b) incubating to hatch and (c) growing the birds for at least 3 weeks after hatch.
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iive 0; Mismatches
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                       Claim 6; Page 37; 45pp; English.
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(GROP-) GROPEP PTY LTD.
(USDA ) US SEC OF AGRIC.
                                                                                                                                                                                                                                                                                                                                                                      N-terminal extension.
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Best Local Similarity
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IGF-1 analogues used pref. (i) lack 1-5 N-terminal amino aicds or (ii) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-1 analogue pref. comprises the N-terminal Given in R51439-49.

IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53.

Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
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IGF-1 analogues used pref. (i) lack 1-5 N-terminal amino aicds or (ii) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced given in R51439-49.
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100.0%; Pre
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Best Local Similarity
Matches 1; Conserv
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IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53.

Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
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100.0%; Pred. No. v.
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Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where
Xaa = any amino acid residue.
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RESULT W56211

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Cyclic derivatives of the peptides also function as anti-inflammatory agents. The peptides can be covalently linked to one another either directly or through a spacer. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus, anti-inflammatory activity. The peptides and compositions have anti-inmune activity. I.e. inhibitory effects against a cellular and humoral immune response, including a response not associated with inflammation. The peptides also inhibit the ability of macrophages and T-cells to adhere to extracellular matrix components and fibronectin, as well as up-regulated fas receptor expression in T-cells. They can be used to inhibit unwanted immune reaction and inflammation.
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                                                                                    Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
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Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where
Xaa = any amino acid residue.
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They are derived from the formulae:
They are din-May, Arg-Glu, Arg-Glu, or Glu-arg-Maa, where
The any amino acid residue

Cyclic derivatives of the peptides also function as anti-inflammatory
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macrophage inhibitory and T-cell inhibitory activity and thus,
anti-inflammatory activity. The peptides and compositions have
anti-inmune activity, i.e. inhibitory effects against a cellular and
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well as up-regulated fas receptor expression in T-cells. They can be used
to inhibit unwanted immune reaction and inflammation.
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Solid phase synthesis; biostatin; cytostatic; tyrosine kinase inhibitor; cellular proliferation inhibition; somatostatin; antitumor.
                                                                                                                                                                                              Peptides W88033-39 are used in the hair composition of the invention. The specification describes a hair composition that comprises at least one oligopeptide having more than two cysteine residues and more than three amino acid residues of the same kind (except cysteine) and a reducing agent. The composition is useful for styling hair to take various shapes and waves safely and effectively.
                                                                                                         New composition for hair - comprises oligopeptide having more than 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Biostatin preparation in high yield by solid synthesis, including closure of disulfide bridge before cleavage from support, useful as antitumor agent -
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                                                                                                                                                               Example 2; Page 6; 17pp; Japanese.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Example 4; Page 23; 33pp; German
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Y51896 standard; peptide; 3 AA.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     99WO-EP06131
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        38WO-EP05306
97JP-0161657
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     16-JUN-2000 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Braum G, Lifferth A,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2000-224663/19
                                                                    WPI; 1999-148442/13
                                                                                                                                                                                                                                                                                                                                                                                               Local Similarity
                                   (LIOY ) LION CORP.
                                                                                                                                                                                                                                                                                                                            3 AA;
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Modified-site
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18-JUN-1997;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                               As a cflu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where Xaa = any amino acid residue.

Xaa = any amino acid residue.

Xaa = any amino acid residue.

Cyclic derivatives of the peptides also function as anti-inflammatory agents. The peptides can be covalently linked to one another either directly or through a spacer. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus, anti-inflammatory activity. The peptides and compositions have anti-inmune activity, i.e. inhibitory effects against a cellular and humoral immune response, including a response not associated with inflammation. The peptides also inhibit the ability of macrophages and T-cells to adhere to extracellular matrix components and fibronectin, as well as up-regulated fas receptor expression in T-cells. They can be used to inhibit unwanted immune reaction and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Gaps
                                                                                                                                                                                                                                                                                                                        Anti-inflammatory peptides and derivatives - used for treating, e.g. arthritis, ulcerative colitis, auto-immune disease, allergy asthma, shock, HIV infection, transplant rejection or Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                    W56171-248 represent anti-inflammatory tripeptides of the invention. They are derived from the formulae:
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100.0%; Pred. No. 0;
iive 0; Mismatches
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                                                                                                                                        97US-0864301.
96US-0025376.
96US-0753141.
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                                                                                                         97WO-IL00295
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Matches 1; Conserv
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                                   WO9809985-A2
                                                                                                         03-SEP-1997;
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                                                                                                                                            28-MAY-1997;
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20-NOV-1996;
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 Synthetic.
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RESULT W88037

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Indels

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This invention describes a novel method for the solid phase (SP)
Synthesis of biostatin (TT 232) (I) which includes closing the disulfide
bridge by oxidation of the completely or partially constructed peptide
while still bonded to the solid phase. The products of the invention have
cytostatic activity and are tyrosine kinase inhibitors. (I) inhibits the
tyrosine kinase activity of various human stomach cancer cell lines and
thus inhibits cellular proliferation. The heptapeptide (I) described in
the invention is a somatostatin analog which shows strong antitumor
cativity in vitro and in vivo. The SP synthesis method gives (I) is more
easily and in markedly higher yield than by the method of EP505680, in
which the cyclization is carried out after cleaving the peptide from the
tresin. The solution method is also a simple synthesis of (I) in high
yield; typically the tert-butyl-protected precursor can be oxidized in
a yield of 70-80%. Y51896-Y51900 represent peptides used in the
synthesis of biostatin TT232 described in the method of the invention.
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3 AA; Sequence

ö Gaps ; 0 Indels Query Match 75.0%; Score 9; DB 21; Length 3; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches 0; Indels

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Search completed: February 5, 2001, 12:01:19 Job time: 366 sec

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POLYPEPTIDE, DNA FRAGMENT ENCODING THE SAME, SAME, AND PROCESS FOR PRODUCING THE SAME, INHIBITION PROCESS, DRUG COMPOSITION AND TREATING USING THE SAME
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         STATE: Virginia
COUNTRY: United States
ZIP: Z2313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/07/791,213D
FILING DATE: 13-NOV-1991
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .,
US-08-079-812-31
US-08-122-510-11
US-08-122-510-12
US-08-122-510-13
US-07-789-913-26
US-08-371-930-4
US-08-372-455-2
US-08-371-585A-1
US-08-321-585A-1
US-08-321-585A-3
US-08-231-1585A-3
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100.0%; Pred. No. 0;
Live 0; Mismatches
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PRIOR APPLICATION: DATA:
PRIOR APPLICATION DATA:
PREDICATION NUMBER: 12-306745
FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: Meuth. Donna M
REGISTRENCE/DOCKET NUMBER: 029650-032
REFERENCE/DOCKET NUMBER: 029650-032
TELECOMMUNICATION INFORMATION:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: MORIGHITA, Hideaki
APPLICANT: MORIGHITA, Hideaki
APPLICANT: NOBJARA, MSABLIDOI
APPLICANT: NOBJARA, MSABLIDOI
TITLE OF INVENTION: POLYPEPTIDE, DNA
TITLE OF INVENTION: INHIBITION PROCES
TITLE OF INVENTION: INHIBITION PROCES
TITLE OF INVENTION: TREATING USING TH
NUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
ADDRESSEE: BUTNS, DOADE, SWECKER &
STREET: P.O. BOX 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ď
                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 23, Application US/07791213D Patent No. 5409895
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Best Local Similarity 100.
Matches 1; Conservative
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STATE: Virginia
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US-07-791-213D-23
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/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6_COMB.pep:*
/cgn2_6/ptodata/2/iaa/PCTUS.COMB.pep:*
/cgn2_6/ptodata/2/iaa/PCTUS.COMB.pep:*
                                     Compugen Ltd
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S-08-354-240A.12
S-08-293-150A-23
S-08-293-150A-39
S-08-65-380-9
S-08-66-390-307
S-08-486-391-307
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S-08-486-391-307
S-08-486-391-307
S-08-249-471-307
S-09-249-451-307
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                     version 4.5
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Maximum Match 100%
Listing first 45 summaries
                                                                                          sw model
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Gapop 10.0 , Gapext 0.5
                       GenCore
Copyright (c) 1993
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                                                                                          protein search, using
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length: 2000000000
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Maximum DB :
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Conservative
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                                                                                                                 COUNTRY: US
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                                                                                                                                                                                                                                                            POLYPEPTIDE, DNA FRAGMENT ENCODING THE SAME, AND ENZYME SAME AND PROCESS FOR PRODUCING THE SAME, AND ENZYME INHIBITION PROCESS, DRUG COMPOSITION AND METHODS OF TREATING USING THE SAME
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Patent No. 5534254
GENERAL INFORMATION, James S.
APPLICANT: Huston, James S.
APPLICANT: Oppermann, Hermann
APPLICANT: Histon, Jewide B.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For
                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Burns, Doane, Swecker & Mathis
STREET: P.O. Box 1404
CITY: ALCANDATION
CITY: D.O. Box 1404
CITY: D.O. Box 1404
CITY: Uriginia
COUNTRY: United States
ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER READABLE FORM:
MEDIUM TYPE: Ploppy disk
COMPUTER IBM PC COMPALIDE
COMPUTER PATENTION DATA:
APPLICATION NUMBER: US/07/791,213D
FILING DATE: 13-NOV-1991
CLASSIFICATION NUMBER: US/07/791,213D
FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: Meuth, Donna M
RECISTRATION NUMBER: 36.607
FELEPHONE: (703) 836-2021
TELEPHONE: (703) 836-2021
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                  Sequence 39, Application US/07791213D Patent No. 5409895
GENERAL INFORMATION:
                                                                                                                                                                                               APPLICANT: MORISHITA, Hideaki
APPLICANT: KARAMORI, TOShinori
APPLICANT: NOBUHARA, Masahirori
TITLE OF INVENTION: POLYPEPTIDE,
TITLE OF INVENTION: SAME AND PROC
TITLE OF INVENTION: INHIBITION PR
TITLE OF INVENTION: TREATING USIN
NUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        LENGTH: 2 amino acids
TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; MOLECULE TYPE: peptide US-07-791-213D-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                 3-07-791-213D-39
Sequence 39, Ap
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US-08-133-804-9
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NUMBER OF SEQUENCE: 11

OURSESPONDER ADDRESS:
ADDRESSEE TREADING Place, 53 State Street
CORRESPONDER ADDRESS:
ADDRESSEE TREADING Place, 53 State Street
STREE: Exchange Place, 53 State Street
STREE: Properties Restreet
STREE: Properties Restreet
STREET: Properties
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0; Indels
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                         Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COMPUTER: TELOPPY disk
COMPUTER: TELOPPY disk
COMPUTER: TENDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: TEM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/293,150A
FILING DATE: 19-AUG-1994
PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/791,213
FILING APPLICATION DATA:
APPLICATION NUMBER: JP 2-306745
FILING DATE: 13-NOV-1990
PRIOR APPLICATION NUMBER: JP 2-306745
FILING DATE: MEUTH, DOING M.
NAME: MEUTH, DOING M.
REGISTERRENCE/DOCKET NUMBER: 36,607
TELECOMMUNICATION: MECHATION:
METERERENCE/DOCKET NUMBER: 029650-049
TELECOMMUNICATION DATA:
METERERENCE/DOCKET NUMBER: 029650-049
                                                               Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Mismatches
                                                                                                                                                                                                                                  Sequence 23, Application US/08293150A Patent No. 5792629 GENERAL INFORMATION:
                                                             :
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                TELEFAX: (703) 836-2021 INFORMATION FOR SEQ ID NO:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   STATE: Virginia
COUNTRY: United States
ZIP: 22313-1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS: LENGTH: 2 amino acids
                                                               Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                , MOLECULE TYPE: peptide US-08-293-150A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
                       Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                  RESULT 6
US-08-293-150A-23
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                                                               Matches
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Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                        0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; LOCATION: 1..2
; OTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"
US-08-461-838-9
                                                                                                                                                                                                                                                                                                                  Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Huston, James S.
APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Departs
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ZIP: 02109
ZIP: 02109
ZIP: 02109
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/NS-DOS
SOSTWARE: PATENTIN Release #1.0, Version #1.25
APPLICATION NUMBER: US/08/461,838
                                                                                                                                                                                                                                                                                                                                                        Mismatches
ATTORNEY/AGENT INFORMATION:
NAME: Sarac, Charles S.
REGISTRATION NUMBER: 30.492
REFERENCE/DOCKET NUMBER: 34506.029
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-831-2100
TELEFAX: 608-831-2106
                                                                                                                                                                                                                                                                                                                75.0%; Score 9;
100.0%; Pred. No
tive 0; Mismat
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US-08-461-838-9
; Sequence 9, Application US/08461838
: Detent No. 5755204
; GENERAL INFORMATION:
                                                                                                             TELEFAX: 608-831-2100
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                TOPOLOGY: linear; MOLECULE TYPE: protein US-08-354-240A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MOLECULE TYPE: protein FEATURE:
                                                                                                                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 1; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 NAME/KEY: Protein
                                                                                                                                                                                                  TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COUNTRY:
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APPLICANT: KANAMORI, TOShinori
APPLICANT: KANAMORI, TOShinori
APPLICANT: NOBIDARA, Masahiro
TITLE OF INVENTION: POLYEPPLIDE, DNA FRAGMENT ENCODING THE
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE SAME, AND ENZYME
TITLE OF INVENTION: TREATING USING THE SAME
TITLE OF INVENTION: TREATING USING THE SAME
NUMBER OF SEQUENCES: 110
CORRESPONDENCE ADDRESS:
STREET: P.O. BOX 1404
CITY: Alexandria
Gaps
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US-08-293-150A-39
; Sequence 39, Application US/08293150A
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Massachusetts
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                                                                                                                       TITLE OF INVENTION: POLYEPTIDE, DNA FRAGMENT ENCODING THE
TITLE OF INVENTION: PARE AND PROCESS FOR PRODUCING THE SAME, AND ENZYME
TITLE OF INVENTION: SAME AND PROCESS, DRUG COMPOSITION AND METHODS OF
TITLE OF INVENTION: TREATING USING THE SAME
NUMBER OF SEQUENCES: 110
CORRESPONDENCE ADDRESS:
ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
STREET: P.O. BOX 1404
CITY: Alexandria
STREET: P.O. BOX 1404
CONNTY: United States
ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
COMPUTER: BROW PC COMPATIBLE
COMPATIBLE READABLE FORM:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                          APPLICANT: MORISHITA, Hideaki
APPLICANT: KANAMORI, Toshinori
APPLICANT: NOBUHARA, Masahiro
TITLE OF INVENTION: POLYPEPIDE, DNA FRAGMENT ENCODING THE
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE SAME, AND ENZYME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           E: Testa, Hurwitz & Thibeault/Patent Department Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 2;
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APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Hing, David B.
TITLE OF INVENTION: Blosynthetic Binding Proteins For INVERTION: Imaging NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/293,150A
FILING DATE: 19-AUG-1994

PLIOR APPLICATION 514

PRIOR APPLICATION NUMBER: US/08/293,150A
FILING DATE: 13-NOV-1990
PRIOR APPLICATION NUMBER: US 07/791,213
FILING DATE: 13-NOV-1990
PRIOR APPLICATION NUMBER: 36,607

FILING DATE: 13-NOV-1990
ATFORNEY/AGENT INFORMATION:

NAME: Meth, Donna M.

REGISTRATION NUMBER: 36,607

REFERENCE/POCKET NUMBER: 36,607

TELEPHONE: (703) 836-6620

TELEPHONE: (703) 836-6620

INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:

LENGTH: 2 main acids

LENGTH: 2 main acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            100.0%; Prec. ...
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TOPOLOGY: linear
MOLECULE TYPE: peptide
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Best Local Similarity
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              US-08-293-150A-39
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                LOCATION: 1..2
CTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"
US-08-461-386-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                APPLICANT: George P. Vlasuk, Patric H. Stanssens, APPLICANT: George P. L. Mensens, Marc J. Lauwereys, APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys, APPLICANT: Yves R. Laroche, Laurent S. Jespers, APPLICANT: Yannick G.J. Gansemans, Matthew Moyle, APPLICANT: Peter W. Bergum
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT TITLE OF INVENTION: PROTEIN
                                                                                                   SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/461,386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 2;
. 0;
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COMPUTER READABLE FORM:
MEDIUM TYPE: -3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,380
FILLING DATE: June 5, 1995
CLASSIFCATION: 530
                  COMPUTER READABLE FORM:
MEDIUM TYPE: "Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ; Sequence 307, Application US/08465380 ; Patent No. 5863894
                                                                                                                                                                                        CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Relley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054,
TELEPOMMUNICATION INFORMATION:
TELEFAX: 617-248-7477
TELEFAX: 617-248-7100
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               75.0%;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        COMPUTER: IBM COMPATIBLE OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                             2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                  STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: 'protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & L)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            NAME/KEY: Protein
LOCATION: 1..2
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           linear
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Best Local Similarity
Matches 1: Conserv
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                                                                                                                                                                             FILING DATE:
02109
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APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: JOLIS H.L. Mensens, Marc J. Lauwereys,
APPLICANT: Yves R. Laroche, Laurent S. Jespers,
APPLICANT: Yves R. Laroche, Laurent S. Jespers,
APPLICANT: Peter W. Bergum
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                 DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             OTHER INFORMATION: xaa in location 2 is an OTHER INFORMATION: amino acid.
                                                                                                                                                                                   Xaa in location 2 is an amino acid.
                                                                                                                                                                                                                                                                                                 75.0%; Score 9; DB 2
100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           COMPUTER PUDDAL

COMPUTER PEADABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 MD

MEDIUM TYPE: Storage

COMPUTER: IBM Compatible

COMPUTER: IBM COMPATIBLE

COMPUTER: IBM COMPATIBLE

OPERATING SYSTEM: IBM P.C. DOS 5.0

CURRENT APPLICATION DATE: US/08/486,399

FILING DATE: June 5, 1995

CLASSIFICATION NUMBER: US/08/486,399

FILING DATE: June 5, 1995

CLASSIFICATION NUMBER: 08/326,110

FILING DATE: OCTOBET 18, 1994

ATTORNEY AGENT INFORMATION:

NAME: BIGGS, SUZANNE L.

RECISTRATION NUMBER: 30,158

REFERENCE/DOCKET NUMBER: 213/270

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION INFORMATION:

TELETAX: (213) 955-0440

TELETAX: 67-3510

INFORMATION FOR SED ID NO: 307:

SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STREET: 633 West Fifth Street STREET: 633 West Fifth Street STREET: Suite 4700 CITY: Los Angeles STATE: California COUNTX: U.S. A ZIP:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; Sequence 307, Application US/08486399
; Patent No. 5866543
                                                                                       TYPE: amino acid
TOPOLOGY: linear
FRACKENT TYPE: internal fragment
         TELEX: 67-3510 INFORMATION FOR SEQ ID NO:
                                                    SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                 1; Conservative
                                                                              2 amino acids
                                                                                                                                                                              OTHER INFORMATION:
COTHER INFORMATION:
US-08-486-397-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
                                                                                                                                                                                                                                                                                                                      Best Local Similarity
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US-08-486-399-307
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                                                                              LENGTH:
                                                                                                                                                                                                                                                                                                                                                                                           2 C 2
                                                                                                                                                                                                                                                                                                      Query Match
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys,
APPLICANT: Yves R. Laroche, Laurent S. Jespers,
APPLICANT: Yannick G.J. Gansemans, Matthew Moyle,
APPLICANT: Peter W. Bergum
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
VITLE OF INVENTION: PROTEIN
WUMBER OF SEQUENCES: 357
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                             Xaa in location 2 is an amino acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Mismatches
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100.0%; Pred. No.
:ive 0; Mismatch
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CLASSIPICATION: 530
CLASSIPICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: October 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/OCKET NUMBER: 213/269
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 90071
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: October 18, 1994
ATTORNEY AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REGISTRATION NUMBER: 213/268
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD Perfect 5.1
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Sequence 307, Application US/08486397 Patent No. 5866542 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                               FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Conservative
                                                                                                                                                                                                                                                                                                                                                                                                           OTHER INFORMATION:
OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 1; Conservat
                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                         US-08-465-380-307
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US-08-486-399-307

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US-08-634-641-307
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CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              0; Indels
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               Length 2;
               DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    ; OTHER INFORMATION: Xaa in location 2 is an ; OTHER INFORMATION: amino acid. US-08-461-965-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              75.0%; Score 9; DB 2;
100.0%; Pred. No. 0;
tive 0; Mismatches
75.0%; Scc. 100.0%; Pred. No. v. 0. Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READABLE FORM:

COMPUTER READABLE FORM:

MEDIUM TYPE: 3.5" Diskette, 1.44 Mb

MEDIUM TYPE: storage

COMPUTER: IBM Compatible

OPERATING SYSTEM: IBM P.C. DOS 5.0

SOFTWARE: Word Perfect 5.1

CURRENT APPLICATION DATA:

APPLICATION NUMBER: US/08/461,965

FILING DATE: Unne 5, 1995

CLASSIFICATION: 5, 1995

CLASSIFICATION DATA:

APPLICATION NUMBER: 08/326,110

FILING DATE: OCTOBE 18, 1994

ATTORNEY/AGENT INFORMATION:

NAME: BIGGS, SUZANNE L.

REGISTRATION NUMBER: 30,138

REGISTRATION NUMBER: 30,138

REGISTRATION NUMBER: 30,138
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TELECOMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                       Sequence 307, Application US/08461965 Patent No. 5872098
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FRAGMENT TYPE: internal fragment
             Query Match 75.0
Best Local Similarity 100.
Matches 1; Conservative
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Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
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US-08-461-965-307
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                                                                 APPLICANT: Vasuk, George P. Vlasuk
APPLICANT: Stanssens, Patrick Eric Hugo
APPLICANT: Stanssens, Joris Hilda Lieven
APPLICANT: Auwerey, Marc Josef
APPLICANT: Laroche, Yves Rene
APPLICANT: Laroche, Yves Rene
APPLICANT: Gansemns, Yannick Georges Jozef
APPLICANT: Georgem, Peter W.
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
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Best Local Similarity 100.0%; Pred. No. 0;
Matches 1; Conservative 0; Mismatches
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COMPUTER: BIBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFFWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/634,641
FILING DATE: APTIL 19, 1996
PRIOR APPLICATION NUMBER: PCT/US95/13231
FILING DATE: October 17, 1995
APPLICATION NUMBER: 08/486,399
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/486,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/465,380
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/466,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: October 18, 1994
ATTONNEY APPORENT THYORMATION:
NAME: PAGGRY THYORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
Sequence 307, Application US/08634641 Patent No. 5955294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME: BIGGS, SUZANNE L. REGISTRATION NUMBER: 30
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Los Angeles
California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ZIP: 90071
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IITLE OF INVENTION: INHIBITORS AND ANTICOAGULANT IITLE OF INVENTION: PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 3;
100.0%; Pred. No. 0;
iive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                             FILING DATE:

PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/809,455
FILING DATE: April 17, 1997
APPLICATION NUMBER: 08/486,399
FILING DATE: Ottober 17, 1995
APPLICATION NUMBER: 08/486,399
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/486,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/46,380
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/46,1965
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/36,110
FILING DATE: October 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANDE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 216/270
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440

INFORMATION FOR SEQ ID NO: 307:
                                                                                                                                                                                                        ZIP: 90071
COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                    COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
FRAGMENT TYPE: internal fragment
                                                                           ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite-4700
CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
TYPE: amino acid
                                     NUMBER OF SEQUENCES: 3
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       OTHER INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ; OTHER INFORMATION:
US-09-249-471-307
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                     U.S.A.
                                                                                                                                                                                   COUNTRY: U
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                                                                                    RESULT 14
US-08-818-253-52
Sequence 52, Application US/08818253
Fatent No. 5998204
GENERAL INFORMATION:
APPLICANT: Tsien, Roger Y.
TITLE OF INVENTION: FLUORESCENT PROFEIN SENSORS
TITLE OF INVENTION: DETECTION OF ANALYTES
NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS: 3
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
STATE: TICA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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TITLE OF INVENTION: NEMATODE-EXTRACTED SERINE PROTEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     MEDIUM TYPE: Diskette
COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FastSEQ for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/818,253
FILING DATE: 14 -MAR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Vlasuk, George Phillip
Stanssens, Patrick Eric Hugo
Messens, Joris Hilda Lieven
Lauwereys, Marc Josef
Laroche, Yves Rene
Jespers, Laurent Stephane
Gansemans, Yannick Georges Jozef
Moyle, Matthew
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100.0%; Pred. No. 0;
tive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ATTORNEY/AGENT INFORMATION:
NAME: Haile, Ph.D., Lisa A.
REGISTRATION NUMBER: 38,347
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          REFERENCE/DOCKET NUMBER: 07
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAX: 619/678-5099
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; MOLECULE TYPE: peptide US-08-818-253-52
                                                                                                                                                                                                                                                                                                                                                                                                                               ZIP: 92037
COMPUTER READABLE FORM:
MEDIUM TYPE: Diskett
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELEPHONE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 RESULT 15
US-09-249-471-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            APPLICANT:
APPLICANT:
APPLICANT:
APPLICANT:
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R-phycocrythrin be proteanine Pl - gor Y protein - human Na+K+-exchanging Glycoprotein compo MHC H2-L antigen - T-cell receptor be T-cell receptor galaminin Bl - weste hypothetical rocket hypothetical prote mabinin II chain vicilin 57K chain cadmium-binding he accetylcholinestera formylglycinamide
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(without alignments)
7.778 Million cell updates/sec
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
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Maximum Match 100%
Listing first 45 summaries

    protein search, using sw model

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Ig H chain V-D-J r T-cell receptor be hypothetical L2 pr Ig heavy chain V r metallothionein is urine glycopeptide L-serine dehydrata leucine-tRNA liga avylase - Kluyvera R-phycoerythrin ga variant surface gl variant surface gl Ig H chain V-D-J r T-cell receptor al gene Cftr protein T-cell receptor be		(Gastroclonium coulteri) (fragment) -Mar-1988 #text_change 23-Mar-1993 tachment sites in R-phycoerythrin.	3 2; Length 3;); ss 0; Indels 0; Gaps 0;) e, P. f the rainbow trout (Oncorhyncus mykiss L/DDBJ PIDN:CAA56466.1; PID:g4379328
PH1602 PH0932 PH0932 SP695 SP622 SP622 A25836 A25836 A37521 C61512 PH1618 PH0803 L57018	ALIGNMENTS	ed alga ision 07 85 bilin at 85182601	Score 9; DB ; Pred. No. 0; 0; Mismatches		ragment) oow trout on 13-Sep of 13-Sep anlysis of 24545 om GB/EMB
00400000000000000000000000000000000000		ter regions	75.0%; 1100.0%; ive		w trout kiss (re nnce_revi Erkell, 349, 199 nnctional mortional mortional
<i>^ ^ ^</i> 0 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8		hrin alpha-1 chain - red algastroclonium coulteri Mar-1988 #sequence_revision V.: A22565 V.: Glazer, A.N. em. 260, 4856-4863, 1985 aracterization of the bilin anumber: A22565; MUID:851826 I mumber: A22565; MUID:851826 type: protein	za t		n-A - rainbow trout (finithy) raphordius mykiss (rain) 1956 #sequence_revision 1049 i Kling, P.; Erkell, L n. 230, 344-349, 1995 n. 230, 344-349, 1995 ininal and functional and functiona
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9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9		RESULT 1 A22565 R-phycocryth C;Species: C C;Date: 07-N C;Accession: R;KlOtz, A.N. J; Biol. Che A;Title: Cha A;Reference A;Accession: A;Molecule t A;Residues:	Query M Best Lo Matches Oy 2	Dþ	RESULT 151049 metalloth C;Species C;Dete: 1 C;Accessil R;Olsson, Eur. J; B A;Referen A;Referen A;Referen A;Referen A;Referen A;Referen A;Referen A;Referen A;Cossil

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C; Species: Datúra innoxia (downy thornapple).
C; Date: 21 May-1990 #text_change 18-Jun-1993
C; Date: 21 May-1990 #sequence_revision 21-May-1990 #text_change 18-Jun-1993
C; Accession: A33882
R; Jackson, P.J; Unkefer, C.J; Doolen, J.A.; Watt, K.; Robinson, N.J.
Proc. Natl. Acad. Sci. U.S.A. 84, 6619-6623, 1997
A; Title: Poly(gamma-qlutamylcysteinyl)glycine: its role in cadmium resistance in plan A; Reference number: A94182; MUID:88016144
A; Accession: A33882
A; Molecule type: protein
A; Residues: 1-5 <JAC>
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C;Species: Plasmodium falciparum
C;Species: Plasmodium falciparum
C;Date: 03-Jun-1993 #sequence_revision 28-Oct-1994 #text_change 09-Jun-2000
C;Accession: B45525 .
R;Wesseling, J.G.; Snijders, P.J.F.; van Someren, P.; Jansen, J.; Smits, M.A.; Schoem Mol. Blochem. Parasital. 35, 167-176, 1989
A;Title: Stage-specific expression and genomic organization of the actin genes of the A;Reference number: A45525; MUID:8936496
R-phycocrythrin gamma-A chain - red alga (Gastroclonium coulteri) (fragment) C;Species: Gastroclonium coulteri C;Decies: Gastroclonium coulteri C;Decies: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993 C;Accession: F22565 A.N.
B:Klotz, A.V.; Glazer, A.N.
J. Biol. Chem. 260, 4856-4863, 1985
A;Title: Characterization of the bilin attachment sites in R-phycocrythrin. A;Reference number: A22565; MUID:85182601
A;Molecule type: proțein
A;Residues: 1-5 <KLO>
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A; Note: the authors translated the codon GAA for residue 3 as
C; Comment: The actin I rank contains
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iive 0; Mismatches
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llarity 100.0%; Pred. No. 0;
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A; Status: preliminary
A; Molecule type: DNA
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555238
pallidiple - assassin bug (fragment)
c;Species: Triatoma pallidipennis (assassin bug)
C;Species: Triatoma pallidipennis (assassin bug)
C;Species: Triatoma pallidipennis (assassin bug)
C;Date: 19-Mar-1997 #sequence_revision 18-Jul-1997 #text_change 19-May-2000
C;Accession: S55238
R;Haendlar, B.; Becker, A.; Noeske-Jungblut, C.; Kraetzschmar, J.; Donner, P.; Schleunin Biochem. J. 307, 465-470, 1995
A;Reference number: S55238; MUID:95251610
A;Reference number: S55238
A;Accession: S55238
A;Accession: S55238
A;Residues: 1-4 <HAE>
                                                                                                            ×.
Ig mu chain V region (clone 13) - human (fragment)
C;Species: Homo sapiens (man)
C;Date: 120-0ct-1994 #sequence_revision 26-Jul-1996 #text_change 17-Mar-1999
C;Accession: $43959
R;Wagner, S.D.; Williams, G.T.; Larson, T.; Neuberger, M.S.; Kitamura, D.; Rajewsky, Nucleic Acids Res. 22, 1389-1393, 1994
A;Title: Antibodies generated from human immunoglobulin miniloci in transgenic mice. A;Reference number: $43956; MuID:94248036
A;Residues: 1-4 <WAG>
C;Keywords: immunoglobulin
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B22565
R-Phycocrythrin alpha-2 chain - red alga (Gastroclonium coulteri) (fragment)
C, Species: Gastroclonium coulteri
C; Dete: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
C; Cate: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
C; Cate: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
C; Cate: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993
R; Molt. A.V.; Glazer, A.N.
J. Biol. Chem. 260, 4856-4863, 1985
A; Title: Characterization of the bilin attachment sites in R-phycoerythrin.
A; Reference number: A22565; MUID:85182601
A; Molecule type: protein
A; Residues: 1-5 < KLO>
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T protein - human (fragment)
C;Species: Homo sapiens (man)
C;Species: Homo sapiens (man)
C;Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 05-Nov-1999
C;Accession: 137263
R;Waeber, G.; Habener, J.F.
Endocrinology 131, 2010-2015, 1992
A;Title: Novel testis germ cell:specific transcript of the CREB gene contains an altc.
A;Reference number: 137263; MUID:93010691
A;Accession: 137263
A;Accession: 137263
A;Molecule type: DNA
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S29881
S29881
S29881
C:Date: Loan's lupus familiaris (dog)
C:Date: 19-Mar-1997 #sequence_revision 24-Jul-1998 #text_change 07-May-1999
C:Accession: S29881
R:Walderhaug, M.O.; Post, R.L.; Saccomani, G.; Leonard, R.T.; Briskin, D.P.
J. Biol. Cham. 260, 3852-3859, 1985
A;Title: Structural relatedness of three ion-transport adenosine triphosphatases arou A;Accession: S29881; MUID:85131201
A;Molecule type: protein
A;Residues: 1-6 CMAL>
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C;Species: Gorilla gorilla)
C;Species: Gorilla gorilla)
C;Species: Gorilla gorilla)
C;Species: 04-oct-1996 #sequence_revision 04-oct-1996 #text_change 21-Jul-2000
C;Accession: 1370a,
R;Queralt, R.; Oliva, R.
Gene 133, 197-204, 1993
A;Title: Identification of conserved potential regulatory sequences of the protamire-A;Reference number: 137013; MUD:94040810
A;Steference number: 137027
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C;Genetics:
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                                                                                                                                                    C; Species: Lumbicus terrestrias (common earthworm)
C; Species: Lumbicus terrestrias (common earthworm)
C; Date: 06-Dec-1996 #sequence_revision 13-Mar-1997 #text_change 13-Mar-1997
C; Accession: 565726
B; C; Accession: 565726
B; C; Accession: 565721; MulD:96176855
A; Accession: 565726
A; Accession: 565
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C: Species: Streptomyces willmorei
C: Date: 20-Apr-2000 #sequence_revision 20-Apr-2000 #text_change 20-Apr-2000
C: Accession: J00355
C: Accession: J00355
B; Tsuda, Y: Okada, Y:; Tanaka, M.; Shigematsu, N.; Hori, Y.; Goto, T.; Hashimoto, M. Chem. Pharm. Bull. 39, 607-611, 1991
A.Title: Structure and synthesis of an immunoactive lipopeptide, WS1279, of microbial A. Riefrence number: J00355; MUID: 91300586
A. Recession: J00355
A. Molecule type: protein
A. Residues: 1-6 < TSUS
A. Molecule type extructure was confirmed by synthesis
C: Keywords: blocked amino end: lipoprotein
F: 1/Binding site: sn-2,3-diacylglycerol (Cys) (covalent) #status experimental
F: 1/Modified site: fatty acylated amino end (Cys) #status experimental
                                                                              S65726 hemoglobin, extracellular, chain b - earthworm (Lumbricus terrestris) (fragment)
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19.Yopyrotein component 16/major fat-globule membrane protein/MFG-E8 homolog - bovine (from 19.Nov-1993 sequence_revision 18-Nov-1994 #text_change 07-Feb-1997 C.Bate: 19-Nov-1993 #sequence_revision 18-Nov-1994 #text_change 07-Feb-1997 C.Accession: H48394
R:Mather, I.H.; Banghart, L.R.; Lane, W.S.
Biochem. Mol. Biol. Int. 29, 545-554, 1993
A;Title: The major fat-globule membrane proteins, bovine components 15/16 and guinea-pig A;Title: The major fat-globule membrane proteins, bovine components 15/16 and guinea-pig A;Title: The major fat-globule membrane proteins, bovine components 15/16 and guinea-pig A;Tetre: H8394
A;Reference number: A48394; MUID:93250576
A;Accession: H48394
A;Actus: preliminary
A;Molecule type: protein
A;Residues: 1-6 cMAT?
A;Experimental source: milk
A;Experimental source: milk
A;Note: sequence extracted from NCBI backbone (NCBIP:131518)
C; Keywords: ATP; heterodimer; hydrolase; ion transport; osmoregulation; phosphoprotein; F; 4/Active site: Asp (aspartylphosphate intermediate) *status predicted
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Search completed: February 5, 2001, 12:02:32 Job time: 289 sec

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us-09-687-267-4.rsp

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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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OM protein - protein search, using sw model

February 5, 2001, 12:06:02; Search time 20.72 Seconds (without alignments) 6.234 Million cell updates/sec Run on:

US-09-687-267-4 12 1 XCXX 4 Title: Perfect score: Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

88757 seqs, 32294092 residues Searched:

88757 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

SwissProt_39:* Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

	Description	; 0	P02729 homo sapien	carci	P05486 conus geogr	couns		_		P42996 scyliorhinu							oryct	P42994 raja clavat				pichia	P17441 pichia jadi		chelyc		couns	lotus	sespan		•		P04558 catostomus	P04559 catostomus
SUMMAKIES	QI	ACT_CARMA	GLUR_HUMAN	CCAP_CARMA	CONO_CONGE	CONO_CONST	DNF1_LOCMI	ISOT_CYPCA	MGMT_BOVIN	OXYA_SCYCA	OXYA_SQUAC	OXYF_SCYCA	OXYT_BUFRE	OXYT_CYPCA	OXYT_EISFO	OXYT_OCTVU	OXYT_RABIT	OXYT_RAJCL	OXYV_SQUAC	RS11_SALTY	SAP_STOVA	TAL1_PICJA	TAL3_PICJA	GLEM_HUMAN	GONZ_CHEPR	RCA_PINPS	CXA1_CONIM	NO40_LOTJA	NO40_SESRO	NO40_SOYBN	RR16_GINBI	TA10_TREME	- 1	UR2B_CATCO
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P04561 cyprinus ca P01147 gillichthys P81022 polyodon sp P35490 scyllorhinu P17776 escherichia P1597 glycine max P15471 conus stria P01520 conus geogr P28878 conus stria P81755 conus texti P81755 conus texti P28525 hordeum vul	S	AA.	date) update)	ireen crab). cea; Malacostraca; eocyemata; Brachyura; Carcinus.	Granjeon E., Webster S.,	.n crab steroidogenesis.";	ID PROTEINS THAT ARE INVOLVED ID ARE UBIQUITOUSLY EXPRESSED	DETERMINED PI OF THIS PROTEIN IS:	FAMILY.	14		NEB3 CRC64;	DB 1; Length 8; 0; :hes 0; Indels 0; Gaps 0;			8 AA.	update) on update)	Craniata; Vertebrata; Euteleostomi; Catarrhini; Hominidae; Homo.
UR2B_CYPCA UR2_FGLEMI UR2_FGLEF UR2_SCYCA YZPY_ECOLI ACT7_SOYBN CXA1_CONST CXA2_CONGE CXA2_CONGE CXA2_CONGE GXA2_CONT CXET_CONT GER1_HORVU	ALIGNMENTS	PRT; 8	<pre>ced) sequence update) annotation updat</pre>	ACIIN (FRAGMENT). Carcinus maenas (Common shore crab) (Green crab) Bukaryota; Metazoa; Arthropoda; Crustacea; Malacc Eumalacostraca; Eucarida; Decapoda; Pleocyemata; Eubrachyura; Portunoidea; Portunidae; Carcinus.	:	An enzyme implicated in	FUNCTION: ACTINS ARE HIGHLY CONSERVED IN VARIOUS TYPES OF CELL MOTILITY AND IN ALL PHYNDROMEN CELLS	YTOPLASMIC	THE ACTIN	ACTINS_1; PARTIAL. ACTINS_2; PARTIAL.	LLINE; FARIT	1424005AB2CAAEB3	Score 9; k; Pred. No. 0; Mismatc			PRT;	sequence	
	* • ·	STANDARD;	(Rel. 34, Created) (Rel. 34, Last seq (Rel. 38, Last ann	r). s (Common sho azoa; Arthroj ; Eucarida; i ortunoidea; i	Somme G., Carpentier G		ACTINS ARE HITTERS OF CENTRAL	R LOCATION: (	6.8, ITS MW IS: 46 KDA. - SIMILARITY: BELONGS TO	05; ACTINS_1	protein.	AA; 976 MW;	75.09 100.0 vative	c		STANDARD;	(Rel. 01, Created) (Rel. 01, Last seq (Rel. 40, Last ann	PTIDE. (Human). :tazoa; Chordata; .heria; Primates;
00000000000000000000000000000000000000		JT 1 SARMA ACT_CARMA	01-0CT-1996 (Re 01-0CT-1996 (Re 15-JUL-1999 (Re	ACIIN (FRAGMENT). Carcinus maenas (Common s) Eukaryota; Metazoa; Arthro Eumalacostraca; Eucarida; Eubrachyura; Portunoidea;	DENCE. haise F.,	bagndassarian D.; "A transaldolase. An en Endocrine 5.23-32(1996)	- FUNCTION: A		6.8, ITS M - SIMILARITY menopo, Ippo	PROSITE; PS00405; PROSITE; PS00432;	Structural prof NON_TER NON_TER	œ	Query Match Best Local Similarity Matches 1; Conser	2 0 2	2 C 2	T 2 HUMAN GLUR_HUMAN P02729.	21-JUL-1986 (Re 21-JUL-1986 (Re 01-OCT-2000 (Re	URINE GLYCOPEPTIDE. Homo sapiens (Human Eukaryota; Metazoa; Mammalia; Eutheria;
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                     Lote C.J., Weiss J.B.;
"Identification in urine of a low-molecular-weight highly polar
"Identification in urine of a low-molecular-weight highly polar
glycopeptide containing cysteinyl-galactose.";
Biochem. J. 123:25P-25P(1971).
-!- FUNCTION: THE IDENTITY OF THE GLYCOPROTEIN FROM WHICH THIS PEPTIDE
IS DERIVED IS UNKNOWN. NO PHYSIOLOGICAL FUNCTION HAS BEEN
ATTRIBUTED. AN ERYTHROCYTE MEMBRANE GLYCOPEPTIDE HAVING A
SIMILAR STRUCTURE HAS ALSO BEEN FOUND.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         'Isolation and identification of a cardioactive peptide from Tenebrio
                                                                                                                                                                                                                   Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         SPECIES-C.MAENAS; TISSUE-PERICARDIAL ORGANS; Stangier J., Hilbiroh C., Beyreuther K., Keller R.; Unusual cardioactive peptide (CCAP) from pericardial organs of the shore crab Carcinus maenas.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SPECIES-M.SEXTA;
MEDLINE-93050243; PubMed-1426284;
MEDLINE-93050243; PubMed-1426284;
Chenng C.C., Loi P.K., Sylwester A.W., Lee T.D., Tublitz N.J.;
Primary structure a cardioactive neuropeptide from the tobacco hawkmoth, Manduca sexta.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    molitor and Spodoptera eridania.";
Bol. Chem. Hoppe-Seyler 374:1065-1074(1993).
-!- FUNCTION: THE EFFECT OF CAP IS BOTH INO- AND CHRONOTROPIC.
-!- FUSSUE SPECIFICITY: STORED IN PERICARDIAL ORGANS AND RELEASED
                                                                                                                                                                                                                   ö
                                                                                                                                                                                                                                                                                                                                                                                               Carcinus maenas (Common shore crab) (Green crab),
Mandicas sexta (Tobacco hawkmoth) (Tobacco hornworm),
Tenebrio molitor (Yellow mealworm), and
Spodoptera eridania (Southern armyworm).
Bukaryotas Metazoa; Arthropoda; Crustacea; Malacostraca;
Eumalacostraca; Eucarida; Decapoda; Plecygemata; Brachyura;
Pubrachyura; Portunoidea; Portunidae; Carcinus.
                                                                                                                                                                                                                   Indels
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MEDILINE-94176032; PubMed-8129851;
FULIVA K., Liao S., Reynolds S.E., Ota R.B., Hackett M.,
Schooley D.A.:
                                                                                                                                                                                           Length 8;
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9 AA; 959 MW; C5A861A9CDD44EB9 CRC64;
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8 AA; 855 MW; C2D87AAlF5BlEB1E CRC64;
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                                                                                                                                                                                                                                                                                                                                                01-0CT-1994 (Rel. 30, Created)
01-0CT-1994 (Rel. 30, Last Sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
CARDIOACTIVE PEFTIDE (CCAP).
                                                                                                                                                                                         75.0%; Score 9; DB 1; ilarity 100.0%; Pred. No. 0; Conservative 0; Mismatches
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SEQUENCE.
MEDLINE=72062338; PubMed=5126885;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Neuropeptide; Amidation.
DISULFID 3 9
                                                                                                                                                                                                                                                                                                                            STANDARD;
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PIR; S27233; S27233.
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Best Local Similarity
                                                                                                                                 Glycoprotein.
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Cruz L.J., de Santos V., Zafaralla G.C., Ramilo C.A., Zeikus R.D.,
Gray W.R., Olivera B.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Cruz L.J., de Santos V., Zafaralla G.C., Ramilo C.A., Zeikus R.D.,
Gray W.R., Olivera B.M.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          "Invertebrate vasopressin/oxytocin homologs. Characterization of peptides from Conus geographus and Conus straitus venoms."; J. Biol. Chem. 262:15821-15824(1987).
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"Peptide toxins: from venomous Conus snails.";
mnu. Rev. Biochem. 57:665-700(1988).
-1- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; A28495; A28495.
INTERPRO; IPRO00981; -.
                                                                                                                                                                                                                                                                                                                                                                       Conus geographus (Geography cone).
Eukaryota; Metazoa; Mollusca; Gastropoda; Caenogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Gastropoda; Caenogastropoda;
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               Length 9;
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D4FC276EB4540059 CRC64;
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(Rel. 09, Last sequence update)
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01-NoV-1988 (Rel. 09, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
ARG-CONOPRESSIN S.
                   DB 1;
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PROSITE: PS00264; NEUROHYPOPHYS_HORM; 1.
Hormone; Amidation.
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Neogastropoda; Conoidea; Conidae; Conus.
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Pred. No.
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9 AA; 1037 MW;
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                 Query Match
Best Local Similarity
Matches 1, Conserva
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Best Local Similarity
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REVIEW

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Rydberg B., Hall J., Karran P.;

"Active site amino acid sequence of the bovine O6-methylguanine-DNA methyltransferase.";

"Nucleic Acids Res. 18:17-21(1990).

-!- FUNCTION: REPAIR OF ALKYLATED GUANINE IN DNA BY STOICHIOMETRICALLY TRANSFRRING THE ALKYL GROUP AT THE O-6 POSITION TO A CYSTEINE RESIDUE IN THE ENZYME. THIS IS A SUICIDE REACTION: THE ENZYME IS IRREVERSIBLY INACTIVATED.

-!- CATALYTIC ACTIVITY: DNA (CONTAINING O6-METHYLGUANINE) + PROTEIN L.CYSTEINE DNA (WITHOUT O6-METHYLGUANINE) + PROTEIN S-METHYL-
                                                                                                                                                                                                                             01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
01-DEC-1995 (Rel. 34, Last annotation update)
METHYLATED-DNA--PROTEIN-CYSTEINE METHYLTRANSFERASE (EC 2.1.1.63) (6-0-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             -!- SIMILARITY: WITH SEGMENTS OF E.COLI ADA AND OGT METHYLTRANSFERASE WHICH ENCOMPASS THE ALKYL-ACCEPTOR RESIDUES.
INTERPRO; IPR001497; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                                                                                                                Cyprinus carpio (Common carp).

Bukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Actinopterygii; Neopterygii; Teleostei; Euteleostei; Ostariophysi;
Cypriniformes; Cyprinidae; Cyprininae; Cyprinus.
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969 MW; 17FF476EB455B04B CRC64;
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. 0;
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01-NOV-1995 (Rel. 32, Last annotation update)
                                9 AA.
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100.0%; Pred. No. 0;
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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                         STANDARD;
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Best Local Similarity
Matches 1; Conserv
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                              ISOT_CYPCA
P42993;
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P29177;
                                                                                                              ISOTOCIN.
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MGMT_BOVIN
                  SOT_CYPCA
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Proux J.P., Miller C.A., Li J.P., Carney R.L., Girardie A.,
Delaage M., Schooley D.A.;
Identification of an arginine vasopressin-like diuretic hormone from
Locusta migratoria.";
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Eukaryota, Metazoa, Arthropoda; Tracheata; Hexapoda; Insecta;
Pterygota, Neoptera; Orthopteroidea; Orthoptera; Caelifera;
Acridomorpha; Acridoidea; Acrididae; Locusta.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Blochem. Blophys. Res. Commun. 149:180-186(1987).
-!- FUNCTION: DIURETIC HORMONE.
-!- SUBUNIT: F2 IS AN ANTIPARALLEL DISULFIDE LINKED DIMER OF
-!- SIMILARITY: BLONGS TO THE VASOPRESSIN/OXTOCIN FAMILY.
PIR: A29477; A39477.
INTERPRO: PIR: PRO00981; -.
PREAM: PF00220; hormone4; 1.
PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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F2).
peptides from Conus geographus and Conus straitus venoms.";
J. Biol. Chem. 262:15821-15824(1987).
                                                                             Gray W.R., Olivera B.M., Cruz L.J.;
"Peptide toxins from venomous Conus snails.";
mnu. Rev. Biochem. 57:665-700(1988).
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; B28495; B28495.
INTERPRO; IPRO00981; -.
PFAM; PF00220; hormone4; 1.
PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
Hormone; Amidation.
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17EB176EB4540050 CRC64;
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56EB176EB451A057 CRC64;
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01-AUG-1990 (Rel. 15, Last Sequence update)
12-DEC-1998 (Rel. 37, Last annotation update)
LOCUPRESSIN (DIURETIC NEUROPEPIIDE F1/F2).
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100.0%; Pred. No. 0;
iive 0; Mismatches
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100.0%; Pred. No. 0;
live 0; Mismatches
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                                                                MEDLINE-89024586; PubMed-3052286;
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Best Local Similarity
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Matches 1; Conserv
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P16339;
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P42997;
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OXYF_SCYCA
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TO 1-NOV-1995 (Rel. 32, Created)

DT 01-NOV-1995 (Rel. 32, Last sequence update)

DT 01-NOV-1995 (Rel. 32, Last annotation update)

C clawryotus, Merazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;

C clasmobranchii; Squalea; Squaloidei; Squalidae; Squalus.

RN FEDLINE-7301727; PubMed=5083097;

RA Acher R., Chauvet J., Chauvet M.-T.;
                                                                                                                               Gaps
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X MEDLINE-95062247; PubMed=7972045;

X Chauvet J., Roulle Y., Chauveau C., Chauvet M.-T., Acher R.;

A Chauvet J., Roulle Y., Chauveau C., Chauvet J., Roulle Y., Chauveau C., Chauvet J., Roulle Y., Chauveau C., Chauvet J., Roulle Y., Experient J., Solation of neurohypophysial hormones in cartilaginous fishes: asvatocin and phasvatocin, two oxytocin-like peptides isolated from the spotted dogfish (Scyliorhinus caniculus).";

Proc. Natl. Acad. Sci. U.S.A. 91:11266-1127(1994).

C -! FUNCTION: DISPLAYS OXYTOCIC ACTIVITY ON RATURY.

INTERRO: IPRO00981; -.

R PFAM: PF00220; hormone4; 1.

R PFAM: PF00220; hormone4; 1.

R HORMONE: Amidation.
                                                                                                                                                                                                                                                                                                                         Scyliorhinus canicula (Spotted dogfish) (Spotted catshark).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchil; Galeomorphil; Galeoidea; Carcharhiniformes;
                                     ALKYL GROUP ACCEPTOR (BY SIMILARITY).
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                                                             325171A720476047 CRC64;
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9 AA; 982 MW; 17EDD76EB444404B CRC64;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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        DNA repair;
        Transferase;
        Methyltransferase.

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        967 MW;
        325171A720476047

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llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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100.0%; Pred. No. 0;
Live 0; Mismatches
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PS00374; MGMT; PARTIAL
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Matches 1; Conserv
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Matches 1; Conserv
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"Phylogeny of the neurohypophysial hormones. Two new active peptides isolated from a cartilaginous fish, Squalus acanthias."; Eur. J. Biochem: 29:12-19(1972).
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyliorhinidae; Scyliorhinus.
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"Special evolution of neurohypophysial hormones in cartilaginous fishes: asvatocin and phasvatocin, two oxytocin-like peptides isolated from the spotted dogfish (Scyliorhinus caniculus)."; Proc. Natl. Acad. Sci. U.S.A. 91:11266-11270(1994).
-i- FUNCTION: DISPLAYS OXYTOCIC ACTIVITY ON RAT UTERUS.
-i- SIMILARITY: 'BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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                                                                                                                                             Acher R., Chauvet J., Chauvet M.-T., Fontaine M.;
"Identification of 2 new neurohypophyseal hormones, valitocin oxytocin) and aspartocin (Ass4-oxytocin) in a selachian fish, spiny dog-fish, Squalus acanthias).",
C. R. Acad. Sci., D., Sci. Nat. 274:313-316(1972).
-! -SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY. INTERPRO; IPRO00981;
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9 AA; 1016 MW; 17EDD76EB44449DB CRC64;
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9 AA; 996 MW; 17F8376EB444404B CRC64;
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100.0%; Pred. No. 0;
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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Pred. No.
                                                                                                                           MEDLINE=72128038; PubMed=4622083;
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100.0%; Pre
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Eisenia foetida (Common brandling worm) (Common dung-worm).
Eukaryota; Metazoa; Annelida; Clitellata; Oligochaeta; Haplotaxida;
Lumbricina; Lumbricidae; Eisenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                MEDLINE-94121660; PubMed-8292046;
Oumi T., Ukena K., Matsushima O., Ikeda T., Fujita T., Minakata H.,
Nomoto K.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      'Annetocin: an oxytocin-related peptide isolated from the earthworm
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Eukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Octopoda;
Incirrata; Octopodidae; Octopus.
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                                                      AMIDATION.
17EB176EB456D04B CRC64;
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01-MAR-1992 (Rel. 21, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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P42998;
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01-NOV-1995
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SPECIES-C.CARPIO; TISSUE-PITUITARY;
Acher R., Chauvet J., Chauvet M.T., Crepy D.;
Characterization of neurohypophyseal hormones from a fresh water bony fish, the carp (Cyprinus carpio). Comparison with hormones from sea fish, the fishs.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                                                                                                                                                                                                                                                                                                                                     Bufo regularis (African toad).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Bufonidae,
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MEDLINE=88225976; PubMed=3371648;
Lane T.F., Sower S.A., Rawauchi H.;
"Arginine vasotocin from the pituitary gland of the lamprey
(Petromyzon marinus): isolation and amino acid sequence.";
Gen. Comp. Endocrinol. 70:152-157(1988).
-!- FUNCTION: ANTIDIURETIC HORMONE.
-!- SIMILARIY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; S01364; B61364.
PIR; S06375; S06375.
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9 AA; 983 MW; 17FF476EA5A6D04B CRC64;
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01-NOV-1991 (Rel. 20, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
                                                01-NOV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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9 AA.
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llarity 100.0%; Pred. No. 0;
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PRT;
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Gaps

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MEDLINE-92270139; PubMed=1589145;

Reich G.;

"A new peptide of the oxytocin/vasopressin family isolated from nerves of the cephalopod Octopus vulgaris.";

Neurosci. Lett. 134:191-194(1992).

-! FUNCTION: HAS A ROLE IN THE NEUROSECRETORY SYSTEM OF THE VENA CAVA.

-! SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.

R PFAM: PF00220; hormone4; 1.

R PFAM: PF00220; hormone4; 1.

R POSITE; PS00264; NEUROHYPOPHYS_HORM; 1.

R POSITE; PS00264; NEUROHYPOPHYS_HORM; 1.

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MEDLINE-93010691; PubMed-1396344;
WEDLINE-93010691; PubMed-1396344;
Waeber G., Habener J.F.;
"Novel testis germ cell-specific transcript of the CREB gene contains an alternatively spliced exon with multiple in-frame stop codons.";
Endocrinology 131:2010-2015(1992).
EMBL; X68994; CAA48780.1; --.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBL_TaxID=9606;
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Conservative 0; Mismatches
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Y PROTEIN (FRAGMENT).
Homo sapiens (Human).
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Copyright (c) 1993 - 2000 Compugen Ltd
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Result

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MEDILINE-81001892; PubMed-7407922;
Dhar R., Chanock R.M., Lai C.J.;
"Nonviral oligonucleotides at the 5' terminus of cytoplasmic influenza viral mrNA deduced from cloned complete genomic sequences.";
cell 21:495-500(1980).
                                                                                                                                                                                                     Viruses; ssRNA positive-strand viruses, no DNA stage; Comoviridae;
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                                                                                                                                                                                                                                                                                                                                                                                   thesis (1992), Biologia Molecular y Virologia Vegetal, CIT-INIA.
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(UDORN/72) HEMAGGLUTININ (SEG 4) CDNA, 3' END (FRAGMENT).
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Viruses; ssRNA negative-strand viruses; Orthomyxoviridae;
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C-TERMINUS OF THE VIRAL REPLICASE (FRAGMENT).
Cherry leaf roll virus.
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MEDLINE-96124520; PubMed-8560786;
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EMBL; 234265; CAA84019.1; -.
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01-NOV-1996 (TrEMBLrel.
01-NOV-1996 (TrEMBLrel.
01-NOV-1998 (TrEMBLrel.
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SEQUENCE 7 AA;
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                                                                                                                                                                                                                                     Nepovirus.
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                                                                                                                                                                                                                                                    Rehm B.H.A., Ertesvag H., Valla S.;
"A new Azotobacter vineland11 mannuronan C-5-epimerase gene (algG) is part of an alg gene Cluster physically organized in a manner similar to that in pseudomonas aeruginosa.";
J. Bacteriol. 178:5884-5889(1996).
BEMBL: X87973; CAA61230.1;
NON TER
SEQUENCE 7 AA; 684 MW; 71B5A5A5D1AEDO CRC64;
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Mammalia; Eutheria; Rodentia; Sciurognathi; Muridae; Murinae; Rattus.
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"New variants of the human and rat nuclear hormone receptor, TR4:
Genomics 35:361-366(1996).
EMBL; U59454; AAB91433.1; -.
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                                                                               Bacteria; Proteobacteria; gamma subdivision; Pseudomonadaceae;
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Yoshikawa I., Makino S., Gao X.M., Xing G.Q., Chuang D.M.,
Detera-Wadleigh S.D.;
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01-JUN-1998 (TrEMBLrel. 06, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
0RPHAN FECEPTOR TR4-NS (FRAGMENT).
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100.0%; Pred. No. 0;
tive 0; Mismatches
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MEDLINE=96299786; PubMed=8661150;
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Best Local Similarity 100.
Matches 1; Conservative
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                                                    Azotobacter vinelandii
  ALGG GENE (FRAGMENT).
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Best Local Similarity
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                                                                                                      Azotobacter.
NCBI_TaxID=354;
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MEDLINE-97442476; PubMed-9295353;
Plummer N.W., Mesurney M.W., Meisler M.H.;
Plummer N.W., Meburney M.W., Meisler M.H.;
Plummer Splicing of the sodium channel SCN8A predicts a truncated two-domain protein in fetal brain and non-neuronal cells.";
J. Biol. Chem. 272:24008-24015(1997).

EMBL: U97673, AAB80916.1; -.
Ionic channel.
NON_TER 1
SEQUENCE 7 AA; 730 MW; 75B72EA2C73772A0 CRC64;
                                                                                                               Human adenovirus type 4.
Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
NCBI_TaxID=28280;
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Actinopterygii; Neopterygii; Teleostei; Buteleostei; Neoteleostei;
Acanthomorpha; Acanthopterygii; Percomorpha; Tetraodontiformes;
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01-NOV-1998 (TIEMBLREI. 08, Last annotation update)
TRUNCATED VOLTAGE-GATED SODIUM CHANNEL ALPHA SUBUNIT (FRAGMENT).
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01-MAY-1999 (TrEMBLrel. 10, Created)
01-MAY-1999 (TrEMBLrel. 10, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
PVI CORE PROTEIN (FRAGMENT).
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01-JAN-1998 (TrEMBLrel. 05,
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STRAIN=KN T96-0620, S-1058, CL 68578;
Crawford-Mikza L.K., Nang R.N., Schnurr D.P.;
Crawford-Mikza L.K., Nang R.N., Schnurr D.P.;
Molecular surveillance of strain variation in adenoviruses causing acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065068; AAD03668.1; -.
EMBL; AF065067; AAD03664.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human adenovirus type 7a.
Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
NCBL_TaxID=85755;
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Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
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Crawford-Miksza L.K., Nang R.N., Schnurr D.P.;
Molecular surveillance of strain variation in adenoviruses
acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAY-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065065; AAD03662.1;
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01-WAY-1999 (TTEMBLrel. 10, Last sequence update)
01-WAY-1999 (TTEMBLrel. 10, Last annotation update)
PVI CORE PROTEIN (FRAGMENT).
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                                                                      (TrEMBLrel. 10,
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Best Local Similarity
Matches 1; Conserv
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SEQUENCE
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TISSUE-PLACENTA;
Lee C.C., Yazdani A., Wehnert M., Bailey J., Couch L., Xiong M.,
Coolbaugh M.I., Chinault C.A., Baldini A., Lindsay E.A., Zhao Z.Y.,
Caskey C.T.H.;
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015900;
01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 07, Last sequence update)
01-AUG-1998 (TrEMBLrel. 07, Last annotation update)
(CLONE XPPIBLA) (FRAMMENT).
(CLONE XPPIBLA) (FRAMMENT).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
(CLONE XP19G12A) (FRACMENT).
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi;
Mammalia, Eutheria, Primates, Catarrhini, Hominidae, Homo.
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
NUCLEAR LIM INTERACTOR (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 AA; 975 MW; 605EA6C5BEA5A2D3 CRC64;
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100.0%; Pred. No. 0;
iive 0; Mismatches
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100.0%; Pred. No. 0;
tive 0; Mismatches
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Hum. Mol. Genet. 0:0-0(0).
EMBL; L32079; AAA73890.1; -.
NON_TER 8 8
SEQUENCE 8 AA, 931 MW; B5D
                                                                                                                                                                                                                                                                                                                                                             Hum. Mol. Genet. 0:0-0(0).
EMBL: L32083; AAA73880.1; -.
NON_TER 1 1 1
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SEQUENCE 8 AA; 975 MW; 6
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TISSUE=PLACENTA;
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0914X6
ID 0914X6
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01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 07, Last sequence update)
01-AUG-1998 (TrEMBLrel. 07, Last annotation update)
(CLONE XP1548A) (FRAGMENT).
Homo sapiens (Human).
Eukaryota: Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
                                                                                                                                                                                   Bacteria; Proteobacteria; gamma subdivision; Enterobacteriaceae;
Escherichia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Hesslinger C., Fairhurst S.A., Sawers G.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
EMBL, AJ001620; CAA04875.1; -.
                                                                                                                                                                                                                                                                                                                                                                          Hesslinger C., Sawers G.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
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                            01-JAN-1998 (TrEMBLrel. 05, Last sequence update) 01-AUG-1998 (TrEMBLrel. 07, Last annotation update) PROPIONATE KINASE (FRAGMENT).
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5. 0;
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tive 0; Mismatches
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(TrEMBLrel. 01, I
(TrEMBLrel. 07, I
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TISSUE-PLACENTA;
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                                                                                                                                                         Escherichia coli
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"Evidence for insufficient chondrocytic differentiation during repair
"Evidence for insufficient chondrocytic differentiation during repair
Mof full-thickness defects of articular cartilage.";
Matrix Biol. 15:39-47(1996).
EMBL; S83371; AAD14433.1; -.
NON_TER 1
SEQUENCE 8 AA; 1028 MW; B859C7272EA77371 CRC64;
                          Drechsler RROW N.A.

Drechsler M., Schumacher V., Friedrich S., Wildhardt G., Giesler S., Schroth A., Bodem J., Royer-Pokora B.;
Schroth A., Bodem J., Royer-Pokora B.;
Genomic structure, alternative transcripts and chromosomal localization of the human LIM domain binding protein 1 gene LDB1/RLI.";
Submitted (JUN-199) to the EMBL/GenBank/DDBJ databases.
BMBL, AJ243097; CAB45408.1; -.
NON_TER 8
SEQUENCE 8 AA; 767 MW; EEGEBDDEB862D5B6 CRC64;
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01-JUL-1997 (TrEMBLrel. 04, Last sequence update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
01-MAY-1999 (TrEMBLrel. 10, Last annotation update)
PRO ALPHA 1 TYPE III COLLAGEN PROTEIN (FRAGMENT).
Oryctolagus cuniculus (Rabbit).
Eukaryota, Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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SEQUENCE FROM N.A.
MEDLINE-96377339; PubMed-8783186;
Metsaranta M., Kujala U.M., Pelliniemi L., Osterman H., Aho H.,
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NCBI_TaxID=9606;
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Search completed: February 5, 2001, 12:05:38 Job time: 289 sec

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Biostatin TT222 sy
Sequence of antiin
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Analgesic and neur
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Synthetic peptide
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Farnesyl-protein t Farnesyl-protein t Fluorophore-labell Farnesyl-protein t Farnesyl-protein t Farnesyl-protein t Farnesyl-protein t

Cyclic platelet ag Cyclic platelet ag Ocular pressure re Antibiotic FR90137 Farnesyl-protein t Farnesyl-protein t Farnesyl-protein t

score:

Title: Perfect s

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Minimum DB Maximum DB

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diabetes.therapy;
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                                                                                                                                                                                                                                                                                                                                                                                                                                            Sequence which corresp. to resides 19-21 of insulin A chain.
                                                                                                                                                                                                                                                                                                                                                                                                                                                              Cystine peptide; insulin activity; hormone; antidiabetic agent.
                                                                                                                                                                                                                                                                                                                                      ALIGNMENTS
                                                      P20217
P30083
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Y51896
P10138
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R15761
Y80840
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R15759
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                                     P10093
P10382
                                                                                                                                                                                                                                                                                                                                                                                    P10301 standard; Protein; 3 AA
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                                                                            80DD-0011868
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(LOSS/) LOSSE G.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Key
Disulfide-bond
\frac{1}{2}
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P10301
Sequence which cor
Binding receptor w
Synthetic peptide
Metal trapping pep
Metal trapping pep
IGF-1 analogue N-t
IGF-1 analogue N-t
IGF-1 analogue N-t
Human interleukin-
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Anti-inflammatory
Anti-inflammatory
Anti-inflammatory
                                                               ; Search time 42.35 Seconds
(without alignments)
3.230 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
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                                                                                                                                                                                                                                                                                   / SIDSI/gcgdata/geneseq/geneseqp/AA1980.DAT:*
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        GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
                                                                                                                                                                                        Total number of hits satisfying chosen parameters:
                                                                                                                                                                       268485 seqs, 34193795 residues
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Maximum Match 100%
Listing first 45 summaries
                                              sw model
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P91660
R34022
R34023
R51440
R51440
W56231
W56231
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Gapop 10.0 , Gapext 0.5
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                                              protein search, using
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Match ]
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Score

Result δ 8 110987654321

Gaps

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The inventors claim synthetic peptides which correspond to sequences occurring in EGF, but excluding EGF. The peptides are angiogenic and have corresp, applications, eg for the healing of wounds and burns. Their relative shortness means that they pose fewer synthesis problems than the entire EGF molecule. They can be admin. singly or association with each other or in association with an angiogenic
                                                                                                                                                                                                                                                           to residues 12-14 of naturally occurring
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Synthetic peptide active in stimulating angiogenesis - has sequences corresponding to amino acid sequences occuring in epidermal growth factor.
     Indels
                                                                                                                                                                                                                                                                                                            Epidermal growth factor; angiogenesis; synthetic peptide
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/note="(aceto amido methyl) NH2-Cys"
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                                                                                                                                                       P91660 standard; protein; 3 AA.
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                                                                                                                                                                                                                                                           Synthetic peptide corresp. to epidermal growth factor (EGF)
                                                                                                                                                                                                                                                                                                                                                                                             /note="H-Gly
                                                                                                                                                                                                                                                                                                                                                                             ./label-OTHER
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     Conservative
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Best Local Similarity
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Misc-difference
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The article for inactivating a toxic material comprises a solid carrier bearing a first receptor which binds the target ligand and a second receptor which degrades the target ligand. This synthetic peptide is a preferred first receptor. The article may be used for covering surfaces to protect or decontaminate the surface. The article is esp. for degrading toxic organic openic openics, besp. organophosphorous cpds., (eg pesticides and nerve poisons, bacteria and viruses, in environmental, chemical, military and industrial settings.
                                                                                      The inventors claim the prepn. of new cysteine peptides with insulin-
                                                                                                    like activity. The new peptides comprise a sequence with corresp. to a central portion of the insulin B chain (esp. B15-27) and a cysteine-conty. peptide with an AA sequence corresp. to a terminal portion of the insulin A chain (esp. A1-7). The new cysteine peptides are antidiabetic agents with appreciable insulin activity (lower than that of insulin itself) and low antigenic activity.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Article for inactivating toxic materials, eg organo:phosporous cpds. comprises solid carrier bearing target ligand binding receptor and ligand-degrading receptor, pref. enzyme
                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                uoxic material; inactivator; organophosphorous cpds.; nerve poison;
pesticide; decontaminant; military.
Cysteine peptide(s) with antidiabetic activity - contg. fragments of insulin A and B chains
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Binding receptor with selectivity for a target ligand, borne by an article for inactivating toxic materials.
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100.0%; Pred. No. 0;
ive 0; Mismatches
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Pred. No. (
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                                                    Claim 1; Page 10; 11pp; German
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100.0%;
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Best Local Similarity
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Best Local Similarity
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RESULT P90982

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peptide (see R31340) with metal trapping activity may be fragment condensation using the peptide fragments given in
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                                                                                                                                                                                                                                                                                                                                                                                                   New acyclic peptide with metal trapping activity - exhibits in vivo metal transporting action and can be used as drug, diagnostic agent or functional material
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in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
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larity 100.0%; Pred. No. 0;
Conservative 0; Mismatches 0; Indels
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                                                /note= "OMe protected"
of cysteine"
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Walton PE;
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(GROP-) GROPEP PTY LTD.
(USDA ) US SEC OF AGRIC.
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Best Local Similarity
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                            Modified-site
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R34022-29.
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   of cysteine"
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                                                                                                                        Electrostatic interaction; metalloprotein; binding site;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Electrostatic interaction; metalloprotein; binding site; protecting group.
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/note= "OMe protected"
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                                                                       Metal trapping peptide intermediate
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             91JP-0120196
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                       19-MAY-1993 (first entry)
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Best Local Similarity
Matches 1; Conserv
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                                                                GF-1 analogues used pref. (1) lack 1-5 N-terminal amino aicds or (ii) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-1 analogue pref. comprises the N-terminal 16F-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. Comprises the N-terminal given in R51450-53.

Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
                                                                                                                                                                                                                                                         Gaps
                        Growth of birds is increased by (a) admin. to the bird, in ovo, insulin-like growth factor (IGF)-1 or -2 or their active analogues; (b) incubating to hatch and (c) growing the birds for at least 3 weeks after hatch.
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in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
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         6; Page 37; 45pp; English.
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                                                                                                                                                                                                                                                                                                                                                 R51440 standard; peptide; 3 AA.
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(GROP-) GROPEP PTY LTD.
(USDA ) US SEC OF AGRIC.
                                                                                                                                                                                   N-terminal extension
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                      3 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                Homo sapiens
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R51440
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weeks after hatch.

IGF-1 analogues used pref. (1) lack 1-5 N-terminal amino aicds or (i1) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-1 analogue pref. comprises the N-terminal given in R51439-49.

IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53.

Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
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IGF-1 analogues used pref. (i) lack 1-5 N-terminal amino aicds or (ii) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced given in R51439-49.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     75.0%; Score 9; DB 15; Length 3; 100.0%; Pred. No. 0; Live 0; Mismatches 0; Indels
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Best Local Similarity
Matches 1; Conserv
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RESULT W19847 W19847;

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W56171-248 represent anti-inflammatory tripeptides of the invention.
They are derived from the formulae:
They can be covaled from the formulae.

Cyclic derivatives of the peptides also function as anti-inflammatory
agents. The peptides can be covalently linked to one another either
Circetly or through a spacer. The peptides and their derivatives have
macrophage inhibitory and T-cell inhibitory activity and thus,
mati-inflammatory activity. The peptides and compositions have
anti-inflammatory activity. The peptides and scapinat a cellular and
humoral immune response, including a response not associated with
T-cells to adhere to extracellular matrix components and fibronectin, as
well as up-requiated fas receptor expression in T-cells. They can be used
to inhibit unwanted immune reaction and inflammation.
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       Gaps
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                                                                                                                                                                                                                                                                                                                           Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
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100.0%; Pred. No. 0;
ive 0; Mismatches
       Mismatches
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                                                                                                                                                                           W56231 standard; peptide; 3 AA
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                                                                                                                                                                                                                                                                                           Anti-inflammatory tripeptide.
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96US-0025376.
96US-0753141.
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                                                                                                                                                                                                                                                      (first entry)
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       Conservative
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nes 1; Conserv
                                                                                                                                                                                                                                                                                                                               Anti-inflammatory;
T-cell inhibitory ?
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   3 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                 WO9809985-A2
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   03-SEP-1996;
20-NOV-1996;
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                                                                                                                                                                                                                                                      20-JUL-1998
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                28-MAY-1997;
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   ij
                                                                                                                                                                                                                                                                                                                                                                                                            Synthetic.
                                         3 C 3
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                                                                                                                                                                                                                  W56231;
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                                                                                                                                      RESULT 10
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     Matches
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IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53.

Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
                                                                                                                                                                                                                                                        Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Interleukin-12; Fc gamma-1; immunosuppressive; autoimmune disease; graft rejection; toxic shock; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              A peptide (W19847) comprises the three N-terminal amino acid residues of human interleukin-12 (IL-12) p40 subunit. It is encoded both by native human p40 cDNA (772096) and by a primer (772095) based on the native sequence. PCR amplification has been used to amplify p40 cDNA. A fusion protein comprising the p40 subunit and human Fc gamma-1 (see also W19853) was expressed in E. coli. This fusion protein has a longer in vivo half-life than native p40 and can be used as an immunosuppressive (e.g. to treat autoimmune diseases or to inhibit graft rejection) or to treat or prevent endotoxin-induced shock.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Fusion protein containing interleukin-12 p40 sub-unit - has increased stability, used to inhibit graft rejection, or treat autoimmune disease and endotoxin-induced shock
                                                                                                                                                                                                                                                      Indels
                                                                                                                                                                                                                  Length 3;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Human interleukin-12 p40 subunit N-terminal peptide.
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                                                                                                                                                                                                               DB 15;
                                                                                                                                                                                                             75.0%; Score 9; DB 1 100.0%; Pred. No. 0; ative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Score 9; I
Pred. No.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  (BETH-) BETH ISRAEL HOSPITAL ASSOC.
                                                                                                                                                                                                                                                                                                                                                                                                                            W19847 standard; Peptide; 3 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Example; Fig 2; 36pp; English.
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100.0%;
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                                                                                                                                                                                                                                                      Conservative
                                                                                                                    N-terminal extension
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                                                                                                                                                                                                                Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
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                                                                                                                                                         3 AA;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Homo sapiens
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W56211;

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RESULT

Synthetic.

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well as up-regulated fas receptor expression in T-cells. They can be used to inhibit unwarted immune reaction and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Cyclic derivatives of the peptides also function as anti-inflammatory agents. The peptides can be covalently linked to one another either directly or through a spacer. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus, anti-inflammatory activity. The peptides and compositions have anti-immune activity, i.e. inhibitory effects against a cellular and humoral immune response, including a response not associated with inflammation. The peptides also inhibit the ability of macrophages and T-cells to adhere to extracellular matrix components and fibronectin, a
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti-inflammatory peptides and derivatives - used for treating, e.g arthritis, ulcerative colitis, auto-immune disease, allergy asthma, shock, HIV infection, transplant rejection or Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                W56171-248 represent anti-inflammatory tripeptides of the invention
                                                         Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                They are derived from the formulae:
Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where
Xaa = any amino acid residue.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                 Eisenbachschwartz M,
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96US-0025376.
96US-0753141.
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                   Anti-inflammatory tripeptide.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Anti-inflammatory tripeptide.
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                        WO9809985-A2.
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20-NOV-1996;
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                                                                                                                                               Synthetic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Sequence
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      W56171-248 represent anti-inflammatory tripeptides of the invention.
They are derived from the formulae:
Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where
Xaa = any amino acid residue.

Cyclic derivatives of the peptides also function as anti-inflammatory
agents. The peptides can be covalently linked to one another either
Circuly or through a spacer. The peptides and their derivatives have
macrophage inhibitory and T-cell inhibitory activity and thus,
anti-inflammatory activity. The peptides and compositions have
anti-inflammatory activity i.e. inhibitory effects against a cellular and
humoral immune response, including a response not associated with
C inflammation. The peptides also inhibit the ability of macrophages and
T-cells to adhere to extracellular matrix components and fibronectin, as
well as up-regulated fas receptor expression in T-cells. They can be used
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                                                                                                                                                                                                        Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
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100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Beserman P, Eisenbachschwartz M,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 6; Page 35; 42pp; English
                                     W56211 standard; peptide; 3 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        W56200 standard; peptide; 3 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (YEDA ) YEDA RES & DEV CO LID
                                                                                                                                                                  Anti-inflammatory tripeptide.
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96US-0753141.
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                                                                                                                         (first entry)
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          3 AA;
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Gaps

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Indels

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Sequence

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W56200;

DXX P

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RESULT W56200 1

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Solid phase synthesis; biostatin; cytostatic; tyrosine kinase inhibitor; cellular proliferation inhibition; somatostatin; antitumor.
                                                                                                                                                                                                      Peptides W88033-39 are used in the hair composition of the invention. The specification describes a hair composition that comprises at least one oligopeptide having more than two cysteine residues and more than three amino acid residues of the same kind (except cysteine) and a reducing agent. The composition is useful for styling hair to take various shapes and waves safely and effectively.
                                                                                                              composition for hair - comprises oligopeptide having more than 2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Biostatin preparation in high yield by solid synthesis, including closure of disulfide bridge before cleavage from support, useful as
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     'note= "Thr(tBu) with C-terminal amide group"
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                                                                                                                                                                                                                                                                                                                                                                                          75.0%; Score 9; DB 2
100.0%; Pred. No. 0;
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biostatin TT232 synthesising peptide 1.
                                                                                                                                 cysteine residues and reducing agent
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               /note= "TFA*Lys(2)"
                                                                                                                                                                     Example 2; Page 6; 17pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; Page 23; 33pp; German.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Y51896 standard; peptide; 3 AA.
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 97JP-0161657.
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                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     /note=
                                                                          WPI; 1999-148442/13
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Best Local Similarity
                                      (LIOY ) LION CORP.
                                                                                                                                                                                                                                                                                                                                   3 AA;
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 18-JUN-1997;
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Matches
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Y51896
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Xaa -[an-Arg, Arg-falu xaa. Xaa-Arg-falu, or Glu-arg-Xaa, where
Xaa = an Arg, Arg-falu xaa. Xaa-Brg-falu, or Glu-arg-Xaa, where
Xaa = any amino acid residue.

Cyclic derivatives of the peptides also function as anti-inflammatory
agents. The peptides can be covalently linked to one another either
directly or through a spacer. The peptides and their derivatives have
macrophage inhibitory and T-cell inhibitory activity and thus,
anti-inflammatory activity. The peptides and compositions have
anti-inflammatory activity. I.e. inhibitory effects against a cellular and
humoral immune activity, i.e. inhibitory effects against a cellular and
humoral immune response, including a response not associated with
inflammation. The peptides also inhibit the ability of macrophages and
r-cells to adhere to extracellular matrix components and fibronectin, as
well as up-regulated fast receptor expression in T-cells. They can be used
to inhibit unwanted immune reaction and inflammation.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                                                                                                                                                                   Anti-inflammatory peptides and derivatives - used for treating, e.g. arthritis, ulcerative colitis, auto-immune disease, allergy asthma, shock, HIV infection, transplant rejection or Alzheimer's disease
                                                                                                                                                                                                                                                                                                                                                                                                                                                W56171-248 represent anti-inflammatory tripeptides of the invention
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                                                                                                                                                                                                                                                               Hirschberg DL;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   75.0%; Score 9; DB 1100.0%; Pred. No. 0; tive 0; Mismatches
                                                                                                                                                                                                                                                               ΣÌ
                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 34; 42pp; English
                                                                                                                                                                                                                                                               Eisenbachschwartz
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         W88037 standard; peptide; 3 AA.
                                                                                                                                                                                                                          (YEDA ) YEDA RES & DEV CO LID
                                                                                                                                              97US-0864301.
96US-0025376.
96US-0753141.
                                                                                                              97WO-IL00295
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Best Local Similarity
Las 1; Conserve
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                                    WO9809985-A2
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Synthetic.

Peptide

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This invention describes a novel method for the solid phase (SP)

Synthesis of biostatin (TT 232) (I) which includes closing the disulfide
bridge by oxidiation of the completely or partially constructed peptide
while still bonded to the solid phase. The products of the invention have
cytostatic activity and are tyrosine kinase inhibitors. (I) inhibits the
tyrosine kinase activity of various human stomach cancer cell lines and
thus inhibits cellular proliferation. The heptapeptide (I) described in
the invention is a somatostatin analog which shows strong antitumor
the invention is a somatostatin analog which shows strong antitumor
cativity in vitro and in vivo. The SP synthesis method gives (I) is more
easily and in markedly higher yield than by the method of EP50580, in
which the cyclization is carried out after cleaving the peptide from the
cell resin. The solution method is also a simple synthesis of (I) in high
yield; typically the tert-butyl-protected precursor can be oxidized in
yield; typically the tert-butyl-protected precursor can be oxidized in
synthesis of biostatin TT232 described in the method of the invention.
              $$$$$$$$$$$$$$$$$$$$
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3 AA; Seguence

ö Gaps ö 0; Indels Query Match 75.0%; Score 9; DB 21; Length 3; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches 0; Indels

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Search completed: February 5, 2001, 12:01:19 Job time: 366 sec

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Sequence 2 Sequence 3 Sequence 1

Sequence 3 Sequence 1 Sequence 2 Sequence 3 Sequence 3

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GENERAL INFORMATION:

APPLICANT: MORISHITA, Hideaki
APPLICANT: KANANORI, TOSHIDOTI
APPLICANT: KANANORI, TOSHIDOTI
APPLICANT: KNOBUHARA, MASAHINO
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE
TITLE OF INVENTION: INHIBITION PROCESS, DRUG COMPOSITION AND
TITLE OF INVENTION: TREATING USING THE SAME
NUMBER OF SOUTENES: 108
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          STATE: ALCASAMILIA
STATE: ALCASAMILIA
COUNTRY: United States
LET: 22313-1404
COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION UNMER: US/07/791,213D
FILING DATE: 13-NOV-1991
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      ;
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US-08-079-812-31
US-08-122-510-11
US-08-122-510-12
US-08-122-510-13
US-07-789-913-26
US-08-371-930-4
US-08-371-930-4
US-08-371-95-2
US-08-371-95-3
US-08-321-585A-3
US-08-321-205A-14
US-08-231-205A-14
US-08-231-205A-14
US-08-231-205A-14
US-08-231-205A-14
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ADDRESSEE: Burns, Doane, Swecker & Mathis STREET: P.O. Box 1404
CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
live 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CLASSITECATION: 433
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JF 2-306745
FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: Meuth, Donna M
REGISTRATION NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 029650-032
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Sequence 23, Application US/07791213D Patent No. 5409895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (703) 836-66.
TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 5
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        1; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Query Match
Best Local Similarity
Matches 1; Conserval
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3Y: linear
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US-07-791-213D-23
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    Sequence 9, Appli
Sequence 23, Appli
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Sequence 9, Appli
Sequence 307, App
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Sequence 39, Appl
                                                                                                                                                                ; Search time 30.91 Seconds (without alignments) 2.324 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         307
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/cgn2_6/ptodata/2/iaa/5B_COMB.pep:*
/cgn2_6/ptodata/2/iaa/6_COMB.pep:*
/cgn2_6/ptodata/2/iaa/pcTUS_COMB.pep:*
/cgn2_6/ptodata/2/iaa/pcTUS_COMB.pep:*
                          GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd
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US-08-451-884-9

US-08-451-884-9

US-08-461-886-9

US-08-465-397-307

US-08-466-397-307

US-08-466-397-307

US-08-466-399-307

US-08-466-399-307

US-08-461-307

US-08-471-307

US-09-249-471-307

US-09-249-461-307

US-07-945-982-8

US-07-947-035-3
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                                                                                                                                                                                                                                                                                                                                                                                                                               174772 seqs, 17957048 residues
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                                                                                                                                                                  2001, 12:01:53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
                                                                                                                       protein search, using sw model
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Gapop 10.0 , Gapext 0.5
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length: 2000000000
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Perfect score:
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RESULT 4
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                                                                                                                                                                                       APPLICANT: MORISHITA, Hideaki
APPLICANT: KANAMORI, TOSHINOTI
APPLICANT: KANAMORI, TOSHINOTI
APPLICANT: NOBUHARA, Masahiro
TITLE OF INVENTION: POLYPEPTIDE, DNA FRAGMENT ENCODING THE
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE SAME, I
TITLE OF INVENTION: INHIBITION PROCESS, DRUG COMPOSITION AND I
TITLE OF INVENTION: TREATING USING THE SAME
NUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
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Patent No. 5534254

GENERAL INFORMATION:
APPLICANT: Usermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Houston, L.
APPLICANT: Houston, Bloaynthetic Binding Proteins For
                                                                                                                                                                                                                                                                                                                                                                             STREET: P.O. Box 1404
CITY: Alexandria
STATE: P.O. Box 1404
COUNTRY: United States
ZJ313-1404
COMPUTER REDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      75.0%; Score .,
100.0%; Pred. No. 0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 2-306745
FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 02950-032
TELECOMMONICATION INFORMATION:
TELEPHONE: (703) 836-6620
TELEPHONE: (703) 836-2021
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           APPLICATION NUMBER: US/07/791,213D FILING DATE: 13-NOV-1991 CLASSIFICATION: 435
                                                                                              RESULT 2
US-07-791-213D-39
; Sequence 39, Application US/07791213D
; Patent No. 5409895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      LENGTH: 2 amino acids TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           , MOLECULE TYPE: peptide US-07-791-213D-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match
Best Local Similarity
                                                                                                                                                                           GENERAL INFORMATION:
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US-08-133-804-9
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                                                                     ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department STREET: Exchange Place, 53 State Street CITY: Boston: STATE: Massachusetts
COUNTRY: USA
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               NAME/KEY: Protein

LOCATION: 1..2

COTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"

US-08-133-804-9
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ZIP: 5317-1914

COMPUTER READABLE FORM:
MEDIUW TYPE: Floppy disk
COMPUTER: IBM PC Compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30

CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/354,240A

FILING DATE: 12-DEC-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                               SOFTWARE: Patentin Release #1.0, Version #1.25 CURRENT APPLICATION DATA: APPLICATION NUMBER: US/08/133,804
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Sequence 12, Application US/08354240A
Patent No. 5670356,
CENERAL INFORMATION:
APPLICANT: Sharf, Bruce A.
APPLICANT: Wood, Keith V.
TITLE OF INVENTION: MODIFIED LUCIFERASE
NUMBER OF SEQUENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCES: 20
CORRESPONDENCES: 3000 CORRESPONDENCES: 3000 CORRESPONDENCES: 3000 Excelsior Drive, Suite 401
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         100.0%; Pred. No. 0; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        75.0%; Score 9;
                                                                                                                                                                                                              COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC COMPATIBLE
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                              CLASSIFICATION: 424
ATTORNEY/AGBAY INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFRENCE/POSKET NUMBER: 2004
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TITLE OF INVENTION: Imaging NUMBER OF SEQUENCES: 11 CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TELEFAX: 617-248-7100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            LENGIH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TYPE: amino acid
STRANDEDNESS: sir
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
Matches 1; Conserv
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                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
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APPLICANT: NOBUHARA, MASAHITO
TITLE OF INVENTION: POLYPEPTIDE, DNA FRAGMENT ENCODING THE
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE SAME, AND ENZYME
TITLE OF INVENTION: INHIBITION PROCESS, DRUG COMPOSITION AND METHODS OF
TITLE OF INVENTION: TREATING USING THE SAME
NUMBER OF SEQUENCES: 110
CORRESPONDENCE ADDRESS:
                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
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                                   Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OPERATING SYSTEM: PC-FOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/293,150A
FILLIG DATE: 19-AUG-1994
CLASSIFICATION: 514
                             Score 9; DB 1;
; Pred. No. 0;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           ADDRESSEE: BJRNS, DOANE, SWECKER & MATHIS STREET: P.O. BOx 1404
CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           PRIOR APPLICATION DATA:
APPLICATION NUMBER: US 07/791,213
FILING DATE: 13-N0V-1990
PRIOR APPLICATION DATA:
APPLICATION NUMBER: JP 2-306745
FILING DATE: 13-N0V-1990
ATTORNEY/AGENT INFORMATION:
NAME: Meuth, Donna M.
REGISTRATION NUMBER: 36,607
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CITY: ALCALL.
STATE: Virginia
COUNTRY: United States
ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM IYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compati
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                75.0%; Score 9;
100.0%; Pred. No
live 0; Mismat
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US-08-293-150A-39
; Sequence 39, Application US/08293150A
                                                                                                                                                                                                                                                                                                                                                     Sequence 23, Application US/08293150A; Patent No. 5792629
        75.0%; Scur
100.0%; Pre
0;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          APPLICANT: KANAMORI, Toshinori
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                REFERENCE/DOCKET NUMBER: 0.
TELECOMMUNICATION INFORMATION
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           INFORMATION FOR SEQ ID NO: SEQUENCE CHARACTERISTICS: LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Conservative
                                                                                      Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; MOLECULE TYPE: peptide US-08-293-150A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity 1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               amino acid
                             Query Match
Best Local Similarity
Matches 1; Conserv
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APPLICANT: MORISH
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STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 LOCATION: 1..2
CTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"
US-08-461-838-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  GENERAL INFORMATION:
APPLICANT: Huston, James S.
APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For TITLE OF INVENTION: Imaging NUMBER OF SEQUENCES: 11
CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,838
                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 1;
100.0%; Pred. No. 0;
ive 0; Mismatches
                       NAME: Sara, Charles S.
REGISTRATION NUMBER: 30,492
REFERENCE/DOCKET NUMBER: 34506.029
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-831-2106
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 9, Application US/08461838
Patent No. 5753204
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NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         34,637
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Conservative
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CITY: Boston
STATE: Massachusetts
                                                                                                                                                                                                                                                                                                                                                  MOLECULE TYPE: protein
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       FILING DATE:
CLASSIFICATION: 424
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                                                                                                                                                                                                                                                                                            TYPE: amino acid
TOPOLOGY: linear
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity
Matches 1; Conserva
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         amino acid
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                                                                                                                                                                                                                                                                                                                                                                                US-08-354-240A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      RESULT 5
US-08-461-838-9
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STATE: Massachusetts
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                        COUNTRY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Matches
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                                                                                                                 POLYPEPTIDE, DNA FRAGMENT ENCODING THE SAME AND PROCESS FOR PRODUCING THE SAME, AND ENZYME THIEITING PROCESS, DRUG COMPOSITION AND METHODS OF TREATING USING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
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APPLICANT: Oppermann, Hermann
APPLICANT: Oppermann, L.
APPLICANT: Houston, L. L.
APPLICANT: Fortan, Hurwitz & Thibeault/Patent Department
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Length 2;
                                                                                                                                                                                                                                                                                                                            COUNTRY: United States

COUNTRY: United States

LIP: 22313-1404

COMPUTER: READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: IBM PC compatible
COMPUTER: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/293,150A
FILING DATE: 19-AUG-1994
CLLASSIFICATION DATA:
APPLICATION NUMBER: US 07/791,213
FILING DATE: 13-NOV-1990
PRIOR APPLICATION NUMBER: US 07/791,213
FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
REGISTRATION NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 36,607
REFINENCE/DOCKET NUMBER: 36,607
REFIRENCE/DOCKET NUMBER: 38-6620
TELEPHONE: (703) 836-6620
                                                                                         TITLE OF INVENTION: POLYPEPTIDE, DNA FRACMENT
TITLE OF INVENTION: SAME AND PROCESS FOR PRODO
TITLE OF INVENTION: INHIBITION PROCESS, DRUG C
TITLE OF INVENTION: TREATING USING THE SAME
NUMBER OF SEQUENCES: 110
CORRESPONDENCE ADDRESS:
ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
STREET: P.O. BOX 1404
CITY: Alexandria
STATE: Virginia
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         75.0%; Score 9; DB 1;
100.0%; Pred. No. 0;
tive 0; Mismatches
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                                            APPLICANT: MORISHITA, Hideaki
APPLICANT: KANAMORI, TOSHINOTI
APPLICANT: NOBUHARA, MASAHIRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 39:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              ; MOLECULE TYPE: peptide US-08-293-150A-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               inear
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Best Local Similarity
Matches 1; Conserv
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                        INFORMATION:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    US-08-461-386-9
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Patent No. 5863894

GENERAL INFORMATION:
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys,
APPLICANT: Tannick G.J. Gansemans, Matthew Moyle,
APPLICANT: Peter W. Bergum
TITLE OF INVENTION: PROTEIN
TITLE OF INVENTION: PROTEIN
TITLE OF INVENTION: PROTEIN

NUMBER OF SEQUENCES:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
COURREPONDER:
CALLEGORIE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ; LOCATION: 1:7
; OTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"
US-08-461-386-9
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURREMY APPLICATION DARA:
APPLICATION NUMBER: US/08/461,386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 2; 100.0%; Pred. No. 0;
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/465,380
FILING DATE: June 5, 1995
                                                                                                                                                                                                    ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REPERENCE/DOCKET NUMBER: 2054
TELECOMMUNICATION:
TELEPHONE: 617-248-7477
TELEFAX: 617-248-7100
                                                                                                                                                                                                                                                                                                                                           TELEFAX: 617-248-7100
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                   TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                             FILING DATE:
CLASSIFICATION: 424
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  NAME/KEY: Protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
APPLICANT: GEOTGE P. Vlasuk, Patric H. Stanssens,
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Joris H.L. Mensens, March S. Jespers,
APPLICANT: Yeas R. Laroche, Laurent S. Jespers,
APPLICANT: Yannick G.J. Gansemans, Matthew Moyle,
APPLICANT: Peter W. Bergum
IITLE OF INVENTION: NEWATODE-EXTRACTED ANTICOAGULANT
IITLE OF INVENTION: PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FEATURE:
OTHER INFORMATION: Xaa in location 2 is an
OTHER INFORMATION: amino acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 75.0%; Score 9; DB 2;
100.0%; Pred. No. 0;
Live 0; Mismatches
                                                                                                                                                                                                                                                                                       ; OTHER INFORMATION: Xaa in location 2 is ; OTHER INFORMATION: amino acid. US-08-486-397-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER SEADABLE FORM:
MEDIUM TYPE: 3.5° Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATE:
APPLICATION WUMBER: US/08/486,399
FILING DATE: June 5, 1995
CLASSIFICATION NUMBER: 08/326,110
FILING DATE: OCCODET 18, 1994
APPLICATION NUMBER: 08/326,110
FILING DATE: OCCODET 18, 1994
ATTORNEY/AGENT INFORMATION:
REGISTRENCE/DOCKET NUMBER: 213/270
TELECOMMUNICATION INFORMATION:
SEQUENCE: 07-3510
INFORMATION FOR SEQ ID NO: 307:
TENERAL OF AMENORMATICE:
TOWN THE OF THE OFFICE O
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        ; Sequence 307, Application US/08486399
; Patent No. 5866543
                                                                                                                                                                                      TOPOLOGY: linear
FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 613 West Fifth Street
STREET: 811te 470
           TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Best Local Similarity 100.
Matches 1; Conservative
                                                                                                                              2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071
                                                                                                                                                        amino acid
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US-08-486-399-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
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Patent No. 5866542

GENERAL INFORMATION

APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys,
APPLICANT: Yves R. Larcche, Laurent S. Jespers,
APPLICANT: Peter W. Bergum

TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT

TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT

TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT

TITLE OF INVENTION: PROTEIN

NUMBER OF SEQUENCES: 357

CORRESPONDENCE ADDRESS:
ADDRESSEE: G33 West Fifth Street

STREET: Suite 4700

CITY: Los Angeles
STREET: California

COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; FEATURE:
    OTHER INFORMATION: Xaa in location 2 is an CTHER INFORMATION: amino acid.
US-08-465-380-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match 75.0%; Score 9; DB 2; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: CALLOCATION

ZIP: 90071

ZIP: 90071

COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 MD
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: WORD PEFFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,397
FILING DATE: June 5, 1995
CLASSIFICATION: 530
PRIOR APPLICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: October 18, 1994
ATTORNEY AGENT INFORMATION:
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: October 18, 1994
ATTORNEY AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 213/268
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 213/269
FELECOMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 US-08-486-397-307
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RESULT 13
US-08-634-641-307
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APPLICANT:
APPLICANT:
APPLICANT:
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                                                                                                                                          APPLICANT:
APPLICANT:
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                                                                                                                                                                                                                                                                            Sequence 307, Application US/08461965

Patent No. 5872098

GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Tyes R. Larcche, Laurent S. Jespers,
APPLICANT: Year R. Larcche, Laurent S. Jespers,
APPLICANT: Yearnick G.J. Gansemans, Matthew Moyle,
APPLICANT: Peter W. Bergum
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
                                                                                                   0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 2;
                                                            Length 2;
                                                          DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   OTHER INFORMATION: Xaa in location 2 is an CTHER INFORMATION: amino acid. US-08-461-965-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             . 0;
. 0;
                                                        75.0%; Score 9; DB 2
100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Query Match 75.0%; Score 9; DB 2
Best Local Similarity 100.0%; Pred. No. 0;
Matches 1; Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       COUNTY. U.S.D.

ZIP: 90071
ZIP: 90071
COMPOTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IEM COMPATIBLE
COMPUTER: MEMORATIBLE
COMPUTER: MEMORATIBLE
COMPUTER: MORE PERFECT 5.1
SOFTWARE: WORD PERFECT 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,965
FILING DATE: US/08/461,965
FILING DATE: OCTOBER 18, 1994
ATTORNEY APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: OCTOBER 18, 1994
ATTORNEY AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
RECESTRATION NUMBER: 30,158
REFERENCE/AGENT NUMBER: 30,158
TELEPHONE: (213) 489-1600
TELEPCOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
TELEPEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FRAGMENT TYPE: internal fragment
                                                                                                   Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             linear
                                                          Query Match
Best Local Similarity
Matches 1; Conserv
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US-08-486-399-307
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Gaps
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                                                                                                                                                                   APPLICANT: Jespers, Laurent Stephane
APPLICANT: Gansemans, Yannick Georges Jozef
APPLICANT: Moyle, Matthew
APPLICANT: Bergum, Peter W.
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
ADDRESSED
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; OTHER INFORMATION: Xaa in location 2 is ; OTHER INFORMATION: amino acid. US-08-634-641-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 2
100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
COMPUTER: 18DM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/634,641
FILING DATE: April 19, 1996
PRICH APPLICATION NUMBER: PCT/US95/13231
FILING DATE: October 17, 1995
APPLICATION NUMBER: 08/466,399
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/466,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/465,380
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/465,380
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: October 18, 1994
ATTORNEY AGENT THORMATION:
NAME: BIGGS, SUZANNE I.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 30,136
                                                                 Vlasuk, George P. Vlasuk
Stanssens, Patrick Eric Hugo
Mensens, Joris Hilda Lieven
Lauwereys, Marc Josef
Laroche, Yves Rene
Jespers, Laurent Stephane
Sequence 307, Application US/08634641
Patent No. 5955294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FRAGMENT TYPE: Internal fragment
                                                                                                                                                                                                                                                                                                                                      STREET: LYON & LYON
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los bre-7
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TELECOMMUNICATION INFORMATION: TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               307:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         TELEPHONE: (213) 489-16(
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: (
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   1; Conservative
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                                                                                                                                                                                                                                                                                                                                                                                                                                 CITY: Los Angeles
STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 1; Conservat
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            amino acid
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                             COUNTRY: U.S.A. ZIP: 90071
                                                GENERAL INFORMATION:
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Length 2;
TITLE OF INVENTION: INHIBITORS AND ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
ADDRESSE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: 633 West Fifth Street
STREET: 633 West Fifth Street
STREET: 631 West Fifth Street
STREET: 105 Angeles
APPLICATION NUMBER: 08/486,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/466,380
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: June 5, 1995
APPLICATION NUMBER: 30,158
REGISTERTEL/OCCKET NUMBER: 216/770
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 3; 100.0%; Pred. No. 0; ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               OTHER INFORMATION: Xaa in location 2 is OTHER INFORMATION: amino acid. US-09-249-471-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Search completed: February 5, 2001, 12:01:54 Job time: 336 sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             TOPOLOGY: linear FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) 489-1600
TELERAX: (213) 955-0440
TELERX: 67-3510
INFORMATION FOR SEQ ID NO: 307
SEQUENCE CHARACIERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Conservative
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Best Local Similarity
Matches 1; Conserva
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                                                                                                                                                                                                                                                     Sequence 52, Application US/08818253
Fatent No. 5998204
GENERAL INFORMATION:
APPLICANT: TSIGN, Actuali
TITLE OF INVENTION: FLUORESCENT PROTEIN SENSORS
TITLE OF INVENTION: FLUORESCENT OF ANALYTES
NUMBER OF SEQUENCES:
ADDRESSEE: Fish & Richardson P.C.
STATE: CA
CITY: La Jolla
STATE: CA
COUNTRY: USA
ZIP: 92037
COMPUTER READABLE FORM:
MEDIUM TYPE: DISKELLE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Indels
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APPLICANT: Bergum, Peter W.
TITLE OF INVENTION: NEMATODE-EXTRACTED SERINE PROTEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Length 2;
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COMPUTER: IBM Compatible
OPERATING SYSTEM: Windows 95
SOFTWARE: FastSEO for Windows Version 2.0b
SOFTWARE: FastSEO for Windows Version 2.0b
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/818,253
FILING DATE: 14-WAR-1997
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 38,347
REGISTRATION NUMBER: 38,347
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 38,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: Vlasuk, George Phillip
APPLICANT: Stanssens, Patrick Eric Hugo
APPLICANT: Stanssens, Joris Hilda Lieven
APPLICANT: Lauwereys, Marc Josef
APPLICANT: Laroche, Yves Rene
APPLICANT: Gaspers, Laurent Stephane
APPLICANT: Gansemans, Yannick Georges Jozef
APPLICANT: Moyle, Matthew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        0; Mismatches
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Patent No. 6040441
GENERAL INFORMATION:
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GenCore version 4.5
Copyright (c) 1993 - 2000 Compugen Ltd.
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OM protein - protein search, using sw model

February 5, 2001, 12:02:32 ; Search time 34.92 Seconds (without alignments) 7.778 Million cell updates/sec Run on:

US-09-687-267-5 12 Title: Perfect score:

1 XXCX 4 Sequence: BLOSUM62 Gapop 10.0 , Gapext 0.5 Scoring table:

195891 seqs, 67900655 residues Searched:

195891 Total number of hits satisfying chosen parameters:

Minimum DB seq length: 0 Maximum DB seq length: 200000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summaries

pir1:* pir2:* pir3:* PIR_66:* 4 3 2 1. .. Database :

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

	Description	phycoerythrin	metallothionein-A	Iq mu chain V reqi	pallidipin - assas	R-phycoerythrin al	R-phycoerythrin ga		•	hemoglobin, extrac	lipopeptide WS1279	R-phycoerythrin be	protamine P1 - gor	- hun	Na+/K+-exchanging	glycoprotein compo	MHC H2-K-k cell su	MHC H2-L antigen -	T-cell receptor be	T-cell receptor ga	laminin B1 - weste	hypothetical TCL3	Ig heavy chain V r	venom heptapeptide	hypothetical prote	mabinlin II chain	vicilin 57K chain	cadmium-binding he	acetylcholinestera	formylglycinamide
SUMMARIES	QI QI	A22565	I51049	S43959	S55238	B22565	F22565	A33882	B45525	S65726	JU0355	C22565	137027	137263	S29881	H48394	I67345	I65546	PT0652	F41946	I49421	179564	PH1408	A58512	808606	S38516	B34818	38	402	A12016
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PH1602 PH0932 156695	PH1407 S59622 XGHUEU	A25836 PC1002 S19288	A37521 C61512 D61512	PH1010 PH00803 PH00934
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## ALIGNMENTS

Riphyocerythrin alpha-1 chain - red alga (Gastroclonium coulteri) (fragment) C;Species: Gastroclonium coulteri) C;Decies: Gastroclonium coulteri) C;Date: 07-Mar-1988 #sequence_revision 07-Mar-1988 #text_change 23-Mar-1993 C;Accession: A22565 R;Klotz, A.V.; Glazer, A.N.
B;Klotz, A.V.; Glazer, A.N.
J;Biol. Chem. 260, 4856-4863, 1985 A;Title: Characterization of the bilin attachment sites in R-phycoerythrin. A;Reference number: A22565; MUID:85182601 A;Molecule type: protein A;Residues: 1-3 <KLO>

ö Gaps ; 0 0; Indels Length 3; DB 2; 75.0%; Score 9; DB 2 100.0%; Pred. No. 0; tive 0; Mismatches 1; Conservative Best_Local Similarity Matches 1; Conserv Query Match

3 C 3 - U ŏ 셤

RESULT I51049

metallothionein-A - rainbow trout (fragment)
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Species: Oncorhynchus mykiss (rainbow trout)
C;Accession: 13-Sep-1996 #text_change 21-Jul-2000
C;Accession: 151049
R;Olsson, P.E.; Kling, P.; Erkell, L.J.; Kille, P.
Bur. J. Biochem. 230, 344-349, 1995
A;Title: Structural and functional analysis of the rainbow trout (Oncorhyncus mykiss)
A;Reference number: 151049
A;Accession: 151049
A;Accession: 151049
A;Accession: 151049
A;Accession: Letter DNA
A;Residues: 1-4 <OLS>
A;Cross-references: EMBL:X80181; NID:91019799; PIDN:CAA56466.1; PID:94379328

Gaps ó 0; Indels Length 4; Ouery Match 75.0%; Score 9; DB 2; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches

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RESULT S43959

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cadmium-binding pentapeptide - downy thornapple
C;Species: Datura innoxia (downy thornapple)
C;Date: 21-May-1990 #sequence_revision 21-May-1990 #text_change 18-Jun-1993
C;Date: 21-May-1990 #sequence_revision 21-May-1990 #text_change 18-Jun-1993
C;Accession: A33882
R;Jackson, P.J.; Unkefer, C.J.; Doolen, J.A.; Watt, K.; Robinson, N.J.
Proc. Natl. Acad. Scl. U.S.A. 84, 6619-6623, 1987
A;Title: Poly(gamma-glutamylcysteinyl)glycine: its role in cadmium resistance in plan A;Reference number: A94182; MUID:88016144
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         C; Species: Plasmodium falciparum
C; Date: 03-Jun-1993 #sequence_revision 28-Oct-1994 #text_change 09-Jun-2000
C; Accession: B45525
R; Wessesling, J.G.; Shijders, P.J.F.; van Someren, P.; Jansen, J.; Smits, M.A.; Schoen Mol. Biochem. Parasitol. 35, 167-176, 1989
A; Title: Stage-specific expression and genomic organization of the actin genes of the A; Reference number: A45525; MUID:89364996
R-phycoerythrin gamma-A chain - red alga (Gastroclonium coulteri) (fragment) C; Species: Gastroclonium coulteri C; Species: Gastroclonium coulteri C; Date: 07-Mar-1988 #text_change 23-Mar-1993 C; Accession: F22565 RKlotz, A.V.; Glazer, A.N. J. Biol. Chem. 260, 4856-4863, 1985 A.Title: Characterization of the billin attachment sites in R-phycoerythrin. A; Reference number: A22565; MUID: 85182601
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A;Note: the authors translated the codon GAA for residue 3 as
C;Comment: The actin I gene contains no introns.
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A, Molecule type: protein
A, Residues: 1-5 <JAC>
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A; Molecule type: DNA
A; Residues: 1-5 <WES>
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pallidipin - assassin bug (fragment)
c;Species Triatoma pallidipennis (assassin bug)
c;Species Triatoma pallidipennis (assassin bug)
c;Species Triatoma pallidipennis (assassin bug)
c;Accession: S55238
R;Haendler, B.; Becker, A.; Noeske-Jungblut, C.; Kraetzschmar, J.; Donner, P.; Schleunin Biochem. J. 307, 465-470, 1995
A;Title: Expression of active recombinant pallidipin, a novel platelet aggregation inhib A;Reference number: S55238; MUID:95251610
A;Accession: S55238
A;Accession: S55238
A;Accession: S55238
A;Residues: 1-4 <ARE>
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$29881

Na+/K-exchanging ATPase (EC 3.6.1.37) alpha chain - dog (fragment)

C;Species: Canis lupus familiaris (dog)

C;Date: 19-War-1997 '*sequence_revision 24-Jul-1998 *text_change 07-May-1999

C;Accession: $29881

R;Walderhaug, M.O.; Post, R.L.; Saccomani, G.; Leonard, R.T.; Briskin, D.P.

J. Biol. Chem. 260, 3852-3859, 1985

A;Title: Structural relatedness of three ion-transport adenosine triphosphatases arou A;Reference number: $29881; MUID:85131201
                                                                                                                                                                                                                         protein PI - gorilla (fragment)

proteines Gorilla gorilla)

C.Date: 04-oct-1996 #sequence_revision 04-oct-1996 #text_change 21-Jul-2000

C.Accession: 137027

R.Queratt, R., Oliva, R.
Gene 133, 197-204, 1993

A.Title: Identification of conserved potential regulatory sequences of the protamine-A.Reference number: 137027

A.Accession: 137027

A.Molecule type: DNA

A.Rolecule type: DNA

A.Rolecule type: DNA

A.Residues: 1-6 <RES>
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137263
C. Protein - human (fragment)
C. Species: Homo sapiens (man)
C. Species: Homo sapiens (man)
C. Date: 12-Aug-1996 #sequence_revision 12-Aug-1996 #text_change 05-Nov-1999
C. Accession: 13763
B. Waber, G.; Habener, J.F.
B. R. Waber, G.; Habener, J.F.
B. A. Feference number: 137263; MUID:93010691
A. Reference number: 137263; MUID:93010691
A. A. Scession: 137263
A. Status: preliminary; translated from GB/EMBL/DDBJ
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C; Genetics:
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                                                                                           Description, extracellular, chain b - earthworm (Lumbricus terrestris) (fragment)
C:Species: Lumbricus terrestris (common earthworm)
C:Species: Lumbricus terrestris (common earthworm)
C:Date: O6-Dec-1996 #sequence_revision 13-Mar-1997
C:Accession: S6726
R:Fushitani, K.; Higashiyama, K.; Asao, M.; Hosokawa, K.
B:Cohim. Biophys. Acta 1292, 273-280, 1996
A:Title: Characterization of the constituent polypeptides of the extracellular hemoglobine Reference number: S65721; MUID:96176855
A:Accession: S6726
A:A
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Cipecies: Orapr-2000 #sequence_revision 20-Apr-2000
Cipecies: Orapr-2000 #sequence_revision 20-Apr-2000
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### Species: Bos primigenius taurus (cattle)
C;Species: Bov-1993 #sequence_revision 18-Nov-1994 #text_change 07-Feb-1997
C;Accession: H48394
R;Mather, I.H.; Banghart, L.R.; Lane, W.S.
Biochem. Mol. Biol. Int. 29, 545-554, 1993
A;Title: The major fat-globule membrane proteins, bovine components 15/16 and guinea-pig
A;Title: The major fat-globule membrane proteins, bovine components 15/16 and guinea-pig
A;Reference number: A48394; MUID:93250576
A;Reference number: H8394
A;Status: preliminary
A;Molecule type: protein
A;Residues: 1-6 <AMIP
A;Residues: 1-6 <AMIP
A;Residues: 1-6 <AMIP
A;Residues: 1-6 <AMIP
A;Residues: 3-6 <AMIP
A;Residues: 3-6 <AMIP
A;Residues: 3-6 <AMIP
A;Residues: 3-6 <AMIP
A;Residues: 3-7 <AMIP
A;Residues: 3-8 <AMIP
A;Residues: 3-10 <AMIP
A;Residues
C; Keywords: ATP; heterodimer; hydrolase; ion transport; osmoregulation; phosphoprotein; F:4/Active site: Asp (aspartylphosphate intermediate) #status predicted
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Non Feb 5 12:13:19 2001
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GenCore version 4.5

Copyright (c) 1993 - 2000 Compugen Ltd.

OM protein - protein search, using sw model

Run on: February 5, 2001, 12:06:02; Search time 20.72 Seconds (without alignments)
6.234 Million cell updates/sec
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Title: US-09-687-267-5 Perfect score: 12 Sequence: 1 XXCX 4

Scoring table: BLOSUM62 Gapext 0.5

Searched: 88757 segs, 32294092 residues

Total number of hits satisfying chosen parameters: 88757

Minimum DB seq length: 0 Maximum DB seq length: 2000000000

Post-processing: Minimum Match 0% Maximum Match 100% Listing first 45 summarles

Database : SwissProt_39:*

Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.

## SUMMARIES

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	Description	P80709	-	_	_	-						P42997												_			-		σ.		_	-	228	P04559 0
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Gaps

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Length 8; 0; Indels

Ouery Match 75.0%; Score 9; DB 1; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches

3 C - C

g G GLUR_HUMAN STANDARD; PRT; 8 AA.
P02729;
1-JUL-1986 (Rel. 01, Created)
21-JUL-1986 (Rel. 01, Last sequence update)
01-OCT-2000 (Rel. 40, Last annotation update)
URINE GLYCOPEPTIDE.
Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
[1]

RESULT 2
GLUR_HUMAN
ID GLUR_HUMAN
AC P02729;
DT 21-JULDT 21-JULDE URINE G
OS HOMO SA
OC BUKATYC
OC MAMMALI
RN [1]

P04561 cyprinus ca P01147 gillichthys P81022 polyodon sp P35490 scyllorhinu P1776 escherichia P15987 glycine max P15987 glycine max P15471 conus stria P01520 conus stria P01520 conus stria P18526 hordeum vul P28525 hordeum vul P28526 hordeum vul			ub). lacostraca; a. prachuira.		n E., Webster S., steroidogenesis.";	PROTEINS THAT ARE INVOLVED ARE UBIQUITOUSLY EXPRESSED	ED PI OF THIS PROTEIN IS:		CRC64;
UR2B_CYPCA UR2_GILMI UR2_POLSP UR2_SCYCA YZPY_ECOLI ACT7_SOYBN CXA_CONST CXA_CONST CXA_CONST CXA_CONST CXA_CONST CXET_CONTE GERT_HORVU	SNMENTS	PRT; 8 AA. ed) sequence update)	ACTIN (FRAGMAT).  Carcinus maenas (Common shore crab) (Green crab).  Bikaryotta, Metazoa, Arthropoda, Crustecea, Malacostraca;  Final arostraca Finantia December 19 consempts	perapora, riecejema Portunidae; Carcinus	, Carpentier G., Granjeon enzyme implicated in crab		CYTOPLASMIC. 2D-GEL THE DETERMINED PI	THE ACTIN FAMILY. PARTIAL. T_LIKE; PARTIAL.	1424005AB2CAAEB3 CF
75.0 75.0 75.0 75.0 75.0 75.0 75.0 12 11 75.0 13 11 75.0 13 11 75.0 13 11 75.0 13 11 75.0	s*.	STANDARD; (Rel. 34, Creat (Rel. 34, Last	13 OUD 1337. ACTIN (FRAGMENT). Carcinus maenas (Common sho Eukaryota, Metazoa, Arthrop	bullaracos raca, bucarina, b Eubrachyura; Portunoidea; P [1] SEQUENCE.	, Soume G., Carpentier an D.; olase. An enzyme implic	Endocrine 5:23-32(1996)!- FUNCTION: ACTINS ARE HIGHLY CONSERVED IN VARIOUS TYPES OF CELL MOTILITY AND IN ALL FURASYOPTIC CELLS.	LOCATION: JS: ON THE IS: 46 KDP	belones 10 1279; 5; ACTINS_1; 2; ACTINS_2; 2; ACTINS_2; 2; ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACTINS_ACT	8 AA; 976 MW;
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P05487;
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CONO_CONST
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                       Lote C.J., Weiss J.B.;
"Identification in urine of a low-molecular-weight highly polar
glycopeptide containing cysteinyl-galactose.";
Biochem. J. 123:25P-25P(1971).
-!- FUNCTION: THE IDENTITY OF THE GLYCOPROTEIN FROM WHICH THIS PEPTIDE
IS DERIVED IS UNKNOWN. NO PHYSIOLOGICAL FUNCTION HAS BEEN
ATTRIBUTED. AN ERYTHROCYTE MEMBRANE GLYCOPEPTIDE HAVING A
SIMILAR STRUCTURE HAS ALSO BEEN FOUND.
PPIR; A03188; XGHUEU.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  "Isolation and identification of a cardioactive peptide from Tenebrio
                                                                                                                                                                                                                                              Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Cheung C.C., Loi P.K., Sylwester A.W., Lee T.D., Tublitz N.J.; "Prinary structure of a cardioactive neuropeptide from the tobacco hawkmoth, Manduca sexta.":
FEBS Lett. 313:165-168(1992).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              molitor and Spodoptera eridania.";
Bol. Chem. Hoppe-Seyler 374:1065-1074(1993).
-!- FUNCTION: THE EFFECT OF CCAP IS BOTH INO- AND CHRONOTROPIC.
-!- TISSUE SPECIFICITY: STORED IN PERICARDIAL ORGANS AND RELEASED
                                                                                                                                                                                                                                              ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                Carcinus maenas (Common shore crab) (Green crab), maduca sexta (Tobacco hawkmoth) (Tobacco hornworm), Tenebrio molitor (Yellow mealworm), and Spodoptera eridania (Southern armyworm).

Bukaryota; Metazoa; Arthropoda; Crustacca; Malacostraca; Eumalacostraca; Bucarida; Decapoda; Plecypemata; Brachyura; Eupirachyura; Portunoidea; Portunoidea; Carcinus.
                                                                                                                                                                                                                                              Indels
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MEDLLINE-94176032; PubMed-8129851;
Furuya K., Liao S., Reynolds S.E., Ota R.B., Hackett M.,
Schooley D.A.
                                                                                                                                                                                                                   Length 8;
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8 AA; 855 MW; C2D87AA1F5B1EB1E CRC64;
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C5A861A9CDD44EB9 CRC64;
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01-0CT-1994 (Rel. 30, Last sequence update)
15-DEC-1998 (Rel. 37, Last annotation update)
CARDIOACTIVE PEPTIDE (CCAP).
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100.0%; Pred. No. 0;
iive 0; Mismatches
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MEDLINE-93050243; PubMed-1426284;
              MEDLINE-72062338; PubMed-5126885;
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PIR; A26363; A26363.
PIR; S27233; S37233.
Deuropeptide; Amidation.
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Best Local Similarity
Matches 1; Conserv
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"Invertebrate vasopressin/oxytocin homologs. Characterization of
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"Peptide toxins from venomous Conus snails.";
Annu. Rev. Biochem. 57:665-700(1988).
-!- SIMILARIY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; A28495; A28495.
PIR; A28495; IPRO00981; -.
PEPAM: PF00220; hormone4; 1.
PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
                                                                                                                                                                                                                                                                                                                                                                                           Conus geographus (Geography cone).
Eukaryota; Metazoa; Mollusca; Gastropoda; Caenogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
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Eukaryota; Metazoa; Mollusca; Gastropoda; Caenogastropoda;
Neogastropoda; Conoidea; Conidae; Conus.
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01-NOV-1988 (Rel. 09, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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Pred. No. 0;
0; Mismatches
                  DB 1;
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Conservative 0;
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1037 MW;
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Best Local Similarity
Matches 1; Conserv
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REVIEW

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Rydberg B., Hall J., Karran P.;
"Active site amino acid sequence of the bovine O6-methylguanine-DNA methyltransferase.";
Nucleic Acids Res. 18:17-21(1990).
-!- FUNCTION: REPAIR OF ALKYLATED GUANINE IN DNA BY STOICHIOMETRICALLY TRANSFERRING THE ALKYL GROUP AT THE O-6 POSITION TO A CYSTEINE RESIDUE IN THE ENZYME. THIS IS A SUICIDE REACTION: THE DIZYME IS
                                                                                                                                                                                                                                                                                                                                                                                           Acher R., Chauvet J., Chauvet M.-T., Crepy D.; "Characterization of neurohypophyseal hormones from a fresh water bony fish, the carp (Cyprinus carpio). Comparison with hormones from sea water bony fishs.";
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     01-DEC-1992 (Rel. 24, Created)
01-DEC-1992 (Rel. 24, Last sequence update)
01-OCT-1996 (Rel. 34, Last annotation update)
METHYLATED-DNA -- PROTEIN-CYSTEINE METHYLARASFERASE (EC 2.1.1.63) (6-O-METHYLGUANINE-DNA METHYLTRANSFERASE) (FRAGMENT).
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CATALYTIC ACTIVITY: DNA (CONTAINING O6-METHYLGUANINE) + PROTEIN

L-CYSTEINE - DNA (WITHOUT O6-METHYLGUANINE) + PROTEIN S-METHYL-
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Cetartiodactyla; Ruminantia; Pecora; Bovoidea;
Bovidae; Bovinae; Bos.
                                                                                                                                                                                                                  Cyprinus carpio (Common carp).

Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Actinopterygii; Neopterygii; Teleostei; Buteleostei; Ostariophysi;
Cypriniformes; Cyprinidae; Cyprinines;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Comp. Biochem. Physiol. 14:245-254(1965).
-!- FUNCTION: ANTIDIURETIC HORMONE.
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY PIR; A61364; A61364.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Length 9;
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9 AA; 969 MW; 17FF476EB455B04B CRC64;
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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100.0%; Pred. No.
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MEDLINE-90174912; PubMed=2308822;
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                                                         STANDARD;
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nes 1; Conserv
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P29177;
                                                         ISOT_CYPCA
P42993;
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SEQUENCE
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  RESULT 7
ISOT_CYPCA
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Proux J.P., Miller C.A., Li J.P., Carney R.L., Girardie A.,
Delaage M., Schooley D.A.;
Identification of an arginine vasopressin-like diuretic hormone
Locusta migratoria.";
Biochem. Biophys. Res. Commun. 149:180-186(1987).
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Eukaryota, Medazoa, Arthropoda; Tracheata, Hexapoda; Insecta,
Pterygota; Neoptera; Orthopteroidea; Orthoptera; Caelifera;
Acridomorpha; Acridoidea; Acrididae; Locusta.
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peptides from Conus geographus and Conus straitus venoms."; J. Biol. Chem. 262:15821-15824(1987).
                                                                                                     MEDLINE-89024586; PubMed-3052286;
Gray W.R., Olivera B.M., Cruz L.J.;
Peptide toxins from venomous Conus snails.";
Annu. Rev. Biochem. 57:665-700(1988).
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR: B28495; B28495.
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-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; A29477; A29477.
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17EB176EB4540050 CRC64;
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01-AUG-1990 (Rel. 15, Last sequence update)
01-BUG-1998 (Rel. 37, Last annotation update)
LOCUPRESSIN (DIURETIC NEUROPEPTIDE F1/F2).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              DB 1;
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llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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ilarity 100.0%;
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9 AA; 1031 MW;
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Best Local Similarity
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"Phylogeny of the neurohypophysial hormones. Two new active peptides isolated from a cartilaginous fish, Squalus acanthias.";
Eur. J. Biochem. 29:12-19(1972).
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"Identification of 2 new neurohypophyseal hormones, valitocin (Val8-oxytocin) and aspartocin (Ash4-oxytocin) in a selachian fish, the spiny dog-fish (Squalus acanthias).";
C. R. Acad. Sci., D, Sci. Nat. 274:313-316(1972).
-!-SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
INTERPRO; IPRO00981;
--PRO0220; hormone4; 1.
PROSITE; PSO0264; NEUROHYPOPHYS_HORM; 1.
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9 AA; 996 MW; 17F8376EB444404B CRC64;
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100.0%; Pred. No. 0;
tive 0; Mismatches
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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9 AA; 1016 MW;
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Best Local Similarity
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DISULFID 1:
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OXYF_SCYCA
ID OXYF_SCYCA
AC P42997;
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Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Chondrichthyes;
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyllorhinidae; Scyllorhinus.
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Eukaryota: Metazoa: Chordata: Craniata: Vertebrata; Chondrichthyes;
Elasmobranchii: Squalea: Squaloidei: Squalidae; Squalus.
                                                                      ALKYL GROUP ACCEPTOR (BY SIMILARITY).
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MEDLINE-95062247; PubMed-7972045;

MEDLINE-95062247; PubMed-7972045;

Chauvet J., Rouille Y., Chauveau C., Chauvet M.-T., Acher R.;

"Special evolution of neurohypophysial hormones in cartilaginous fishes: asvatocin and phasvatocin, two oxytocin-like peptides isolated from the spotted dogfish (Scyllorhinus caniculus).";

Proc. Natl. Acad. Sci. U.S.A. 91:11266-11270(1994).

-i- FINCTION: DISPLAYS OXYTOCIC ACTIVITY ON RAT UTERUS.

-i- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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9 AA; 982 MW; 17EDD76EB444404B CRC64;
                                                                                                             325171A720476047 CRC64
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
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100.0%; Pred. No. 0;
iive 0; Mismatches
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01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
ASPARTOCIN (ASPARGTOCIN)
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PROSITE; PS00374; MGMT; PARTIAL.

DNA repair; Transferase; Methyltransferase.

NON_TER 1 1 ALKYL GROUP ACI
NON_TER 9 9 SEQUENCE 9 AA; 967 MW; 325171A720476047
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PROSITE: PS00264; NEUROHYPOPHYS_HORM; 1.
HORMONE; Amidation.
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Acher R., Chauvet J., Chauvet M.-T.;
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P42996;
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ID OXYA_SQUAC
AC P42999;
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Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Chondrichthyes,
Elasmobranchii; Galeomorphii; Galeoidea; Carcharhiniformes;
Scyllorhinidae; Scyllorhinus.
                                                                                                                                                                                                                                                                                                                      MEDLINE-95062247; PubMed-7972045;
MEDLINE-95062247; PubMed-7972045;
MEDLINE-95062247; PubMed-7972045;
MEDLINE-95062247; PubMed-7972045;

"Special evolution of neurohypophysial hormones in cartilaginous fishes: asvatocin and phasvatocin, two oxytocin-like peptides isolated from the spotted dogfish (Scyllorhinus caniculus).";
proc. Natl. Acad. Sci. 0.5A. 91:11266-11270(1994).
-1- FUNCTION: DISPLASS OXYTOCIC ACTIVITY ON RAT UTERUS.
-1- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
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17EDD76EB44449DB CRC64;
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01-NoV-1995 (Rel. 32, Created)
01-NOV-1995 (Rel. 32, Last sequence update)
01-NOV-1995 (Rel. 32, Last annotation update)
PHASYATOCIN.
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Hormone; Amidation.
DISULFID 1.
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P42998;
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Acher R., Chauvet J., Chauvet M.-T., Crepy D.;
"Characterization of neurohypophyseal hormones from a fresh water bony fish, the carp (Cyprinus carpio). Comparison with hormones from sea
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                     MEDINE-96099313; PURMORITEMENT LODE;
Chauvet J., Michel G., Ouedraogo Y., Chou J., Chait B.T., Acher R.;
"A new neurohypophysial peptide, seritocin ([Ser5,Ile8]-oxytocin),
identified in a dryness-resistant African toad, Bufo regularis.";
Int. J. Pept. Protein Res. 45:482-487(1995).
-!-FUNCTION: DEVOID OF OXYTOCIC ACTIVITY.
-!- SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
INTERPRO, IPRO00981; -.
PROSITE; PS00220; hormone4; 1.
PROSITE; PS00264; NEUROHYPOPHYS_HORM; FALSE_NEG.
DISULEID
                                                                                                                    Bufo regularis (African toad).
Eukaryota, Metazoa, Chordata, Craniata, Vertebrata, Euteleostomi,
Amphibia, Batrachia, Anura, Neobatrachia, Bufonoidea, Bufonidae,
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MEDLINE—88225976; PubMed=3314648;
Lane T.F., Sover S.A., Kawauchi H.;
Lane T.F., Sover S.A., Kawauchi H.;
Arginine vasotocin from the pituitary gland of the lamprey (Petromyzon marinus): isolation and amino acid sequence.";
Gen. Comp. Endocrinol. 70:152-157 (1988).
-!- FUNCTION: ANTIDIURETIC HORMONE.
-!- FUNCTION: ANTIDIURETIC HORMONE.
PIR; 801364, B61364.
PIR; S06375; S06375.
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9 AA; 983 MW; 17FF476EA5A6D04B CRC64;
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9 AA.
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llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
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Comp. Biochem. Physiol. 14:245-254(1965)
                                                                                                                                                                                                                                          TISSUE=PITUITARY NEUROINTERMEDIATE LOBE;
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PROSITE; PS00264; NEUROHYPOPHYS_HORM; 1.
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Biochem Biophys. Res. Commun. 198:393-399(1994).
-! FUNCTION: POTENTIATES SPONTANEOUS CONTRACTIONS OF THE GUT AND ALSO PULSATORY CONTRACTIONS AND BLADDER-SHAKING MOVEMENT OF THE NEPHRIDIA. MAY BE INVOLVED IN OSMOREGULATION OF THE ANIMAL THROUGH NEPHRIDIAL FUNCTION.
-! SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
PIR; PC2021; PC2021.
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Eukaryota; Metazoa; Annelida; Clitellata; Oligochaeta; Haplotaxida;
Lumbricina; Lumbricidae; Eisenia.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                "Annetocin: an oxytocin-related peptide isolated from the earthworm
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Oumi T., Ukena K., Matsushima O., Ikeda T., Fujita T., Minakata H., Nomoto K.;
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Eukaryota; Metazoa; Mollusca; Cephalopoda; Coleoidea; Octopoda;
Incirrata; Octopodidae; Octopus.
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tive 0; Mismatches
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TISSUE=NERVE ENDINGS;
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MEDLINE-92270139; PubMed-1589145;
Reich G.;
The we peptide of the oxytocin/vasopressin family isolated from nerves of the cephaloped octopus vulgaris.";
Neurosci. Lett. 134:191-194(1992).
-! FUNCTION: HAS A ROLE IN THE NEUROSECRETORY SYSTEM OF THE VENA CANA.
-! SIMILARITY: BELONGS TO THE VASOPRESSIN/OXYTOCIN FAMILY.
R PFAM: PF00220; hormone4; 1.
R PFAM: PF00220; hormone4; 1.
R PRAM: PF00220; hormone4; 1.
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Q47063 escherichia
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Q9987 homo sapien
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Q9409 monodelphis
G183040 monodelphis
G183040 acrococcus
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G69473 human herpe
O12096 caprine art
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047063 escherichia
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MEDLINE=93010691; PubMed=1396344;
Waeber G., Habener J.F.;
Waeber Itestis germ cell-specific transcript of the CREB gene contains an alternatively spliced exon with multiple in-frame stop codons.";
Endocrinology 131:2010-2015(1992).
EMBL; X68994; CAA48780.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Homo sapiens (Human).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
NCBI_TaxID=9606;
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SECUENCE FROM N.A.
MEDLINE-81001992; bubMed=7407922;
Dhar R., Chancok R.M., Lai C.J.;
"Nonviral oligonucleotides at the 5' terminus of cytoplasmic influenza viral mRNA deduced from cloned complete genomic sequences.";
Call 21:495-500(1980).
EMBL; M25045; AAA43202.1; -.
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0.10V-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
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c-TEMAINOS OF THE VIRAL REPLICASE (FRAGMENT).
c-TEMAINOS OF THE VIRAL REPLICASE (FRAGMENT).
Viruses; ssRNA positive-strand viruses, no DNA stage; Comoviridae;
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"Long, nearly identical untranslated sequences at the 3' terminal regions of the genomic RNAs of cherry leafroll virus (walnut strain).";
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(UDORN/72) HEMAGGLOTININ (SEG 4) CDNA, 3' END (FRAGMENT).
Influenza A virus.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Viruses; ssRNA negative-strand viruses; Orthomyxoviridae;
Influenza virus A and B group.
NCBL_TaxID=11320;
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100.0%; Pred. No. 0;
iive 0; Mismatches
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larity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
                 7 AA.
                   PRT;
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EMBL; 234265; CAA84019.1; -.
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                   PRELIMINARY;
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NCBI_TaxID=12615;
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                                                                                                                                                               MEDLINE=96427318; PubMed=8830682; Rehm B.H.A., Ertesvag H., Valla S.; Rehm B.H.A., Ertesvag H., Valla S.; "A new Azotobacter vinelandii mannuronan C-5-epimerase gene (algG) is part of an alg gene cluster physically organized in a manner similar to that in Pseudomonas aeruginosa."; J. Barteriol. 178:5884-5889[1996). EMBL; X87973; CAA61230.1; -.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Rattus norvegicus (Rat).
Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Rođentia; Sciurognathi; Muridae; Murinae; Rattus.
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Yoshikawa T., Dupont B.R., Leach R.J., Detera-Wadleigh S.D.;
"New variants of the human and rat nuclear hormone receptor, TR4:
expression and chromosomal localization of the human gene.";
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                                                      Bacteria; Proteobacteria; gamma subdivision; Pseudomonadaceae;
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Last annotation update)
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100.0%; Pred. No. 0;
tive 0; Mismatches
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                                  Azotobacter vinelandii
ALGG GENE (FRAGMENT)
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Best Local Similarity
Matches 1; Conserv
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                                                   Human adenovirus type 4.
Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
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Bukaryota: Metazoa; Chordata: Craniata: Vertebrata: Euteleostomi;
Actinopterygil; Neopterygil; Teleostel; Euteleostel; Neoteleostel;
Acanthomorpha; Acanthopterygil; Percomorpha; Tetraodontiformes;
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01-JAN-1998 (TrEMBLrel. 05, Last sequence update)
01-NOV-1998 (TrEMBLrel. 08, Last annotation update)
TRUNCATED VOLTAGE-GATED SODIUM CHANNEL ALPHA SUBUNIT (FRAGMENT).
SCN8A.
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Plummer N. W., McBurney M.W., Meisler M.H.;
Palternative splicing of the sodium channel SCN8A predicts
two-domain protein in fetal brain and non-neuronal cells.";
J. Biol. Chem. 272:24008-24015(1997).
EMBL; U97673; AAB80916.1; -.
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7 AA; 980 MW; 7B5EA414140322A0 CRC64;
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01-MAY-1999 (TrEMBLrel. 10, 01-MAY-1999 (TrEMBLrel. 10, 01-MAY-1999 (TrEMBLrel. 10, PVI CORE PROTEIN (FRAGMENT).
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01-JAN-1998 (TrEMBLrel. 05,
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                                                                                                       Human adenovirus type 7.
Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
NCBI_TaxID=10519;
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Viruses; dsDNA viruses, no RNA stage; Adenoviridae; Mastadenovirus.
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Crawford-Miksza L.K., Nang R.N., Schnurr D.P.;
Molecular surveillance of strain variation in adenoviruses acute respiratory disease, AV 4 and AV 7a.";
Submitted (MAX-1998) to the EMBL/GenBank/DDBJ databases.
EMBL; AF065065; AAD03662.1;
                                                                                                                                                                                                                                                                 75.0%; Score 9; DB 12; Length 7; 100.0%; Pred. No. 0; O. 100els ive 0; Mismatches 0; Indels
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Matches 1; Conserv
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Lee C.C., Yazdani A., Wehnert M., Bailey J., Couch L., Xiong M.,
Coolbadyh M.I., Chinault C.A., Baldini A., Lindsay E.A., Zhao Z.Y.,
Caskey C.T.H.;
Hum. Mol. Genet: 0:0-0(0).
EMBL; L32083; AAA73880.1; -.
NON_TER 8 8
SEQUENCE 8 AA, 975 MW; 605EA6C5BEA5A2D3 CRC64;
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(CLONE XP19G12A) (FRAGMENT).

Homo sapiens (Human).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;

Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Homo sapiens (Human).
Bukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi;
Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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Mammalia; Eutheria; Primates; Catarrhini; Hominidae; Homo.
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01-NOV-1999 (TrEMBLrel. 12, Last sequence update)
01-NOV-1999 (TrEMBLrel. 12, Last annotation update)
NUCLEAR LIM INTERACTOR (FRAGMENT).
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100.0%; Pred. No. 0;
tive 0; Mismatches
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100.0%; Pred. No. 0;
iive 0; Mismatches
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                                                                                                                                                                                                                                         Homo sapiens (Human).
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Best Local Similarity
Matches 1; Conserv
                                                                                          SEQUENCE FROM N.A.
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Q15900;
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Q15900
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               01-NOV-1996 (TIEMBLIEL. 01, Created)
01-NOV-1996 (TIEMBLIEL. 01, Last sequence update)
01-NOV-1998 (TIEMBLIEL. 07, Last annotation update)
01-NOV-1998 (TIEMBLIEL. 07, Last annotation update)
(CLONE XP15H8A) (FRAGMENT).
Homo sapiens (Human).
Homo sapiens (Human).
Manalia: Metazoa; Chordata; Craniata; Vertebrata; Buteleostomi;
Manmalia: Butheria; Primates; Catarrhini; Hominidae; Homo.
                                                                       Bacteria; Proteobacteria; gamma subdivision; Enterobacterlaceae;
Escherichia.
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Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
EMBL; AJ001620; CAA04875.1; -
                                                                                                                                                           Hesslinger C., Sawers G.;
Submitted (SEP-1997) to the EMBL/GenBank/DDBJ databases.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 4; Length 8; 100.0%; Pred. No. 0; tive 0; Mismatches 0; Indels
                                                                                                                                                                                                                                                                                                                       Length 8;
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01-NOV-1996 (TrEMBLrel. 01, Created)
01-NOV-1996 (TrEMBLrel. 01, Last sequence update)
01-AUG-1998 (TrEMBLrel. 07, Last annotation update)
           01-JAN-1998 (TIEMBLEEL. 05, Last sequence update) 01-AUG-1998 (TIEMBLEEL. 07, Last annotation update) PROPIONATE KINASE (FRAGMENT).
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             8 AA; 1068 MW; 0315A37EAB5B0763 CRC64;
                                                                                                                                                                                                                                                                                SEQUENCE 8 AA; 1000 MW; 3A505EB044140DC4 CRC64;
                                                                                                                                                                                                                                                                                                                       75.0%; Score 9; DB 2; 100.0%; Pred. No. 0; vative 0; Mismatches
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EMBL: L32069; AAA73878.1; -.

NON_TER 8

SEQUENCE 8 AA; 1068 MW; C
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Best Local Similarity
Matches 1; Conserv
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                                                                                                    NCBI_TaxID=562;
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"Evidence for insufficient chondrocytic differentiation during repair
"Evidence for insufficient chondrocytic differentiation during repair
of full-thickness defects of articular cartilage.";
Matrix Biol. 15:38-47(1996).
EMBL; S83371; AAD14433.1; -
NON_TER
SEQUENCE 8 AA; 1028 MW; B859C7272EA77371 CRC64;
                                                  Gaps
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01-JUL-1997 (TIEMBLE 1. 04, Last sequence update)
01-MAY-1999 (TIEMBLE 1. 10, Last annotation update)
PRO ALPHA 1 TYPE III COLLAGEN PROTEIN (FRAGMENT).
Cryctolagus cuniculus (Rabbit).

Eukaryota; Metazoa; Chordata; Craniata; Vertebrata; Euteleostomi; Mammalia; Eutheria; Lagomorpha; Leporidae; Oryctolagus.
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SEQUENCE FROM N.A.
MEDLINE-96377339; PubMed-8783186;
Metsaranta M., Kujala U.M., Pelliniemi L., Osterman H., Aho H.,
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Search completed: February 5, 2001, 12:05:38 Job time: 289 sec

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Anti-inflammatory Peptide used in th Biostatin TT232 sy

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W56211 W56200

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Thioredoxin active Cyclic platelet ag Cyclic platelet ag Ocular pressure re Antibiotic FR90137 Farnesyl-protein t Farnesyl-protein t

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Farnesyl-protein Farnesyl-protein Farnesyl-protein Farnesyl-protein Fluorophore-label ; 0

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Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             The inventors claim synthetic peptides which correspond to sequences occurring in EGF, but excluding EGF. The peptides are angiogenic and have corresp, applications, eg for the healing of wounds and burns. Their relative shortness means that they pose fewer synthesis problems than the entire EGF molecule. They can be admin. Singly or association with each other or in association with an angiogenic
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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Pred. No. 0;
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                                                                                                                                                      protein; 3 AA.
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epidermal growth factor (EGF)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                The article for inactivating a toxic material comprises a solid carrier bearing a first receptor which binds the target ligand and a second receptor which degrades the target ligand. This synthetic peptide is a preferred first receptor. The article may be used for covering surfaces to protect or decontaminate the surface. The article is esp. adequating toxic organic cyds, esp. organophosphorous cyds., (eg pesticides and nerve poisons, bacteria and viruses, in environmental, chemical, military and industrial settings.
                                                                                                  like activity. The new peptides comprise a sequence with corresp. to a central portion of the insulin B chain (esp. B15-27) and a cysteine-conty. peptide with an AA sequence corresp. to a terminal portion of the insulin A chain (esp. A1-7). The new cysteine peptides are antidiabetic agents with appreciable insulin activity (lower than that of insulin itself) and low antigenic activity.
                                                                                    The inventors claim the prepn. of new cysteine peptides with insulin-
                                                                                                                                                                                                                                                                                                          Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Toxic material; inactivator; organophosphorous cpds.; nerve poison; pesticide; decontaminant; military.
Cysteine peptide(s) with antidiabetic activity - contg. fragments of insulin A and B chains
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100.0%; Pred. No. v.
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                                                     Claim 1; Page 10; 11pp; German
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100.0%;
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Matches

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The acyclic peptide (see R31340) with metal trapping activity may be produced by fragment condensation using the peptide fragments given in R34022-29.
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                                                                                                                                                                                                                New acyclic peptide with metal trapping activity - exhibits in vivo metal transporting action and can be used as drug, diagnostic agent or functional material
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Insulin-like growth factor; IGF-1; IGF-2; bird; chicken; egg;
in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
                                                                                                                                                                                                                                                                                                                                                                                                                         Indels
                                                                                                                                                                                                                                                                                                                                                                                               DB 14; Length 3; 0;
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                                                                                                                                                                                                                                                                                                                                                                                             75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         McMurtry JP,
                       /note- "OMe protected"
                                                                                                                                                             KK.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               R51439 standard; peptide; 3 AA.
                                                                                                                                                                                                                                                                        Claim 3; Page 2; 9pp; Japanese
                                                                                                                                                            (ICHI-) ICHIKAWA GOSEI KAGAKU
                                                                                                       91JP-0120196.
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(GROP-) GROPEP PTY LID.
(USDA ) US SEC OF AGRIC.
                                                                                                                                                                                       WPI; 1993-021449/03.
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                                                                                                                                                                                                                                                                                                                                                                                                          Local Similarity
Les 1; Conserv
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           Modified-site
                                                  JP04346999-A.
                                                                                                        24-MAY-1991;
                                                                                                                                  24-MAY-1991;
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Walton PE;
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                                                                              02-DEC-1992
                                                                                                                                                                                                                                                                                                                                                       Seguence
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R51439
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Gaps
                                                                                                                                                                                     /note= "Cys(Acm); Acm= acetamidomethyl-protected SH
   of cysteine"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  /note= "Cys(Acm); Acm= acetamidomethyl-protected SH
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0
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Electrostatic interaction; metalloprotein; binding site; protecting group.
                                                                Electrostatic interaction; metalloprotein; binding site;
                                                                                                                                                         note= "Boc protected"
                                                                                                                                                                                                                               /note= "OMe protected"
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Metal trapping peptide intermediate.
                                    Metal trapping peptide intermediate.
                                                                                                                                   Location/Qualifiers
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Location/Qualifiers
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                                                                                                                                                                                                                                                                                                                                                                   KK.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Claim 3; Page 2; 9pp; Japanese
                                                                                                                                                                                                                                                                                                                                                                   (ICHI-) ICHIKAWA GOSEI KAGAKU
                                                                                                                                                                                                                                                                                                                                        91JP-0120196.
         19-MAY-1993 (first entry)
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Best Local Similarity
Matches 1; Conserv
                                                                              protecting group
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           3 AA;
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Modified-site
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Modified-site
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                                                                                                        Synthetic.
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Gaps

Gaps

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Indels

Claim 6;

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weeks after hatch.

IGF-1 analogues used pref. (i) lack 1-5 N-terminal amino aicds or (ii) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-1 analogue pref. comprises the N-terminal given in R51439:49.

IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53.

Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
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IGF-1 analogues used pref. (i) lack 1-5 N-terminal amino aicds or (ii) have Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced given in R51439-49.
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in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
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                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
iive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         R51441 standard; peptide; 3 AA.
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(GROP-) GROPEP PIX LID.
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Best Local Similarity
Matches 1; Conserva
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                                                                                                                                                                                                                                                                                                       3 AA;
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                                                                                                                                            TGF-1 analogues used pref. (i) lack 1-5 N-terminal amino aicds or 1GF-1 analogues used pref. (i) lave Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-1 analogue pref. comprises the N-terminal given in R51439-49.

IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53.

Esp. Long R3 IGF-1, given in R51451 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Gaps
                                                                  Growth of birds is increased by (a) admin. to the bird, in ovo, insulin-11ke growth factor (IGF)-1 or -2 or their active analogues; (b) incubating to hatch and (c) growing the birds for at least 3 weeks after hatch.
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in ovo; growth; promotion; increase; long R3 IGF-1; LR3 IGF-1.
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100.0%; Pred. No. 0;
ive 0; Mismatches
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                          Page 37; 45pp; English
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(GROP-) GROPEP PTY LTD.
(USDA ) US SEC OF AGRIC.
                                                                                                                                                                                                                                                                                                                                                                                             N-terminal extension.
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                    3 AA;
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Anti-inflammatory peptides and derivatives - used for treating, e.g. arthritis, ulcerative colitis, auto-immune disease, allergy asthma, shock, HIV infection, transplant rejection or Alzheimer's disease
                                                                                                                                                                                                                                                                                        Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
     Mismatches
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                                                                                                                                                      W56231 standard; peptide; 3 AA.
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96US-0753141.
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Best Local Similarity 100.
Matches 1; Conservative
     Conservative
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                                                                                                                                                                                                                        20-JUL-1998
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20-NOV-1996;
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     Matches
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IGF-2 analogues used pref. have Glu(2) or Glu(3) absent or replaced by Gly, Gln, Leu, Arg or Lys, partic. Gly and the Thr normally adjacent to Glu can be replaced by Arg or Gly. The IGF-2 analogue pref. comprises the N-terminal given in R51450-53. Esp. Long R3 IGF-1, given in R51454 is used. It is the full human IGF-1 sequence with Arg replacing Glu(3) and a 13 amino acid
                                                                                                                                                                                                                          Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Interleukin-12; Fc gamma-1; immunosuppressive; autoimmune disease; graft rejection; toxic shock; therapy.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                A peptide (W19847) comprises the three N-terminal amino acid residues of human interleukin-12 (IL-12) p40 subunit. It is encoded both by native human p40 cDNA (T72095) and by a primer (T72095) based on the native sequence. PCR amplification has been used to amplify p40 cDNA. A fusion protein comprising the p40 subunit and human Fc gamma-1 (see also W19853) was expressed in E. coll. This fusion protein has a longer in vivo half-life than native p40 and can be used as an immunosuppressive (e.g. to treat autoimmune diseases or to inhibit graft rejection) or to treat or prevent endotoxin-induced shock.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Fusion protein containing interleukin-12 p40 sub-unit - has increased stability, used to inhibit graft rejection, or treat autoimmune disease and endotoxin-induced shock
                                                                                                                                                                                                                        Indels
                                                                                                                                                                                       Length 3;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Human interleukin-12 p40 subunit N-terminal peptide.
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                                                                                                                                                                                       DB 15;
                                                                                                                                                                                    75.0%; Score 9; DB 1
100.0%; Pred. No. 0;
ive 0; Mismatches
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                                                                                                                                                                                                                                                                                                                                                                          W19847 standard; Peptide; 3 AA.
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                                                                                                                                                                                                                      Conservative
                                                                                                    N-terminal extension.
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N-PSDB; T72095-96.
                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                     3 AA;
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W19847
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NAME OF COLOR OF STREET STREET

Hirschberg DL;

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                                                                                                                           Again, authorized of the peptides also function as anti-inflammatory agents. The peptides can be covalently linked to one another either directly or through a spacer. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus, anti-inflammatory activity. The peptides and compositions have anti-inflamma activity, i.e. inhibitory effects against a cellular and humoral immune response, including a response not associated with inflammation. The peptides also inhibit the ability of macrophages and T-cells to adhere to extracellular matrix components and fibronectin, as well as up-regulated fas receptor expression in T-cells. They can be used to inhibit unwanted immune reaction and inflammation.
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W56171-248 represent anti-inflammatory tripeptides of the invention. They are derived from the formulae:
Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where
any amino acid residue.
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100.0%; Pre
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Length 3;

DB 18;

75.0%; Score 9; I 100.0%; Pred. No.

Query Match Best Local Similarity

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Anti-inflammatory tripeptide.
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les 1; Conserv
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03-SEP-1996;
20-NOV-1996;
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                                                                                                                                                                                                                                                                 03-SEP-1997;
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                                                                                                                                                                                                                           12-MAR-1998
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Xaa Glu-Axq, Axq-Glu-Xaa, Xaa-Axq-Glu, or Glu-arg-Xaa, where
Xaa any amino acid residue.

Xaa any amino acid residue.

Cyclic derivatives of the peptides also function as anti-inflammatory agents. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus anti-inflammatory activity. The peptides and compositions have anti-inflammation activity. The peptides and compositions have anti-inmune activity. In the peptides and compositions have inflammation. The peptides also inhibit the ability of macrophages inflammation. The peptides also inhibit the ability of macrophages and T-cells to adhere to extracellular matrix components and fibronectin, as well as up-regulated fas receptor expression in T-cells. They can be used to inhibit unwanted immune reaction and inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Hirschberg DL;
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                                    W56211 standard; peptide; 3 AA.
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                                                                                                                                                             Anti-inflammatory tripeptide.
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96US-0753141.
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                                                                            W56211;
RESULT 11
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Matches
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They are derived from the formulae:

Xaa-Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where

Xaa = any amino acid residue.

Cyclic derivatives of the peptides also function as anti-inflammatory
agents. The peptides can be covalently linked to one another either
agents. The peptides can be covalently linked to one another either
alizectly or through a spacer. The peptides and their derivatives have
macrophage inhibitory and T-cell inhibitory activity and thus,
anti-inflammatory activity. The peptides and compositions have
anti-inmune activity, i.e. inhibitory effects against a cellular and
humoral immune response, including a response not associated with
inflammation. The peptides also inhibit the ability of macrophages and
T-cells to adhere to extracellular matrix components and fibronectin, as
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inhibit unwanted immune reaction and inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     W56171-248 represent anti-inflammatory tripeptides of the invention
Anti-inflammatory; macrophage inhibitory activity; fibronectin; T-cell inhibitory activity; adherence; extracellular matrix; up-regulation; fas receptor expression; inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Claim 5; Page 34; 42pp; English.
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96US-0025376.
96US-0753141.
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us-09-687-267-6.rag

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Solid phase synthesis; biostatin; cytostatic; tyrosine kinase inhibitor; cellular proliferation inhibition; somatostatin; antitumor.
                                                                                                                                                                                        Peptides W88033-39 are used in the hair composition of the invention. The specification describes a hair composition that comprises at least one oligopeptide having more than two cysteine residues and more than three amino acid residues of the same kind (except cysteine) and a reducing agent. The composition is useful for styling hair to take various shapes and waves safely and effectively.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Biostatin preparation in high yield by solid synthesis, including closure of disulfide bridge before cleavage from support, useful as antitumor agent -
                                                                                                      comprises oligopeptide having more than
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                /note= "Thr(tBu) with C-terminal amide group"
                                                                                                                                                                                                                                                                                                                                                                      Length 3;
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                                                                                                                                                                                                                                                                                                                                                                    DB 20;
                                                                                                                                                                                                                                                                                                                                                                                                          Mismatches
                                                                                                                                                                                                                                                                                                                                                                      Score 9; 1
Pred. No.
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                                                                                                                         cysteine residues and reducing agent
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                                                                                                                                                        Example 2; Page 6; 17pp; Japanese.
                                                                                                                                                                                                                                                                                                                                                          75.0%; Scor.
100.0%; Pre
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97JP-0161657.
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                                                                                                        composition for hair
                                                                                                                                                                                                                                                                                                                                                                                                        Conservative
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                                                                    WPI; 1999-148442/13
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Best Local Similarity
Matches 1; Conserv
                                  (LIOY ) LION CORP.
                                                                                                                                                                                                                                                                                                                    3 AA;
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Modified-site
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18-JUN-1997;
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Xaa Glu-Arg, Arg-Glu-Xaa, Xaa-Arg-Glu, or Glu-arg-Xaa, where
Xaa = any amino acid residue.

Cyclic derivatives of the peptides also function as anti-inflammatory agents. The peptides can be covalently linked to one another either directly or through a spacer. The peptides and their derivatives have macrophage inhibitory and T-cell inhibitory activity and thus, anti-inflammatory activity. The peptides and compositions have anti-inflammatory activity. The peptides and compositions have humoral immune activity, i.e. inhibitory effects against a cellular and humoral immune response, including a response not associated with inflammation. The peptides also inhibit the ability of macrophages and reals a calcular matrix components and fibronectin, as used to a constitution of the peptides also inhibit and inflammation.
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                                                                                                                                                                                                                                                                                                                                                                                      Claim 3; Page 34; 42pp; English.
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                                                                                                                                    97US-0864301.
96US-0025376.
96US-0753141.
                                                                                                        97WO-IL00295.
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Matches 1; Conserv
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20-NOV-1996;
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                                                                    12-MAR-1998
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Synthetic
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Sequence

Query Match

δ g Peptide

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This invention describes a novel method for the solid phase (SP)

Synthesis of biostatin (TT 232) (I) which includes closing the disulfide

Dridge by oxidation of the completely or partially constructed peptide

while still bonded to the solid phase. The products of the invention have

Cytostatic activity and are tyrosine kinase inhibitors. (I) inhibits the

tyrosine kinase activity of various human stomach cancer cell lines and

thus inhibits cellular proliferation. The perpapentide (I) described in

the invention is a somatostatin analog which shows strong antitumor

ceasily and in markedly higher yield than by the method of EP50580, in

which the cyclization is carried out after cleaving the peptide from the

cesin. The solution method is also a simple synthesis of (I) in high

cresin. The solution method is also a simple synthesis of (I) in high

cycled; typically the terributyl protected precursor can be oxidized in

a yield of 70-80%. Y51896-Y51900 represent peptides used in the

synthesis of biostatin TT232 described in the method of the invention.
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3 AA; Sequence

ö 0; Gaps Query Match 75.0%; Score 9; DB 21; Length 3; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches 0; Indels

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Search completed: February 5, 2001, 12:01:19 Job time: 366 sec

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         POLYPEPTIDE, DNA FRAGMENT ENCODING THE SAME AND PROCESS FOR PRODUCING THE SAME, INHIBITION PROCESS, DRUG COMPOSITION AND 17REATING USING THE SAME
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OPPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PATENTIN Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
FILING DATE: 13-NOV-1991
CLASSIFICATION: 435
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US-08-079-812-31
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US-08-371-930-4
US-08-372-455-8
US-08-321-585A-1
US-08-321-585A-1
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US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-321-585A-3
US-08-331-585A-3
US-08-331-585A-3
US-08-331-585A-3
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US-08-231-205A-14
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Mismatches
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100.0%; Pred. No
:ive 0; Mismat
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APPLICATION NUMBER: JP 2-306745
FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: Meuth, Donna M
REGISTRATION NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 029650-03
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence 23, Application US/07791213D Patent No. 5409895
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     APPLICANT: MORISHITA, Hideaki
APPLICANT: KANAMORI, TOShinori
APPLICANT: NOBJHARA, MASAHIRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ZIP: 22313-14v.4
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
naururer: IBM PC compatible
naururer: IBM PC compatible
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      APPLICANT: NOBJARRA, MASABILIO
TITLE OF INVENTION: POLYPEPTI
TITLE OF INVENTION: SAME AND
TITLE OF INVENTION: INHIBITIO
TITLE OF INVENTION: TREATING
NUMBER OF SEQUENCES: 108
CORRESPONDENCE ADDRESS:
ADDRESSEE: BUIRS, DOANE, SW
STREET: P.O. BOX 1404
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 TELEPHONE: (703) 836-6620
TELEFAX: (703) 836-2021
INFORMATION FOR SEQ ID NO: 23:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          CITY: Alexandria
STATE: Virginia
COUNTRY: United States
ZIP: 22313-1404
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Matches 1; Conservative
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GENERAL INFORMATION:
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US-08-354-240A-12

US-08-293-150A-29

US-08-293-150A-39

US-08-465-380-307

US-08-465-397-307

US-08-465-397-307

US-08-461-307-307

US-08-461-307

US-08-634-641-307

US-08-249-471-307

US-09-249-471-307

US-09-249-461-307

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Maximum Match 1008
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Sequence:
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                                                                                                                                                                                                                 APPLICANT: MORISHITA, Hideaki
APPLICANT: KANAMORI, Toshinori
APPLICANT: NOBUHARA, Masahiro
TITLE OF INVENTION: POLYPEPTIDE, DNA FRAGMENT ENCODING THE
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE SAME, AND ENZYME
TITLE OF INVENTION: INHIBITION PROCESS, DRUG COMPOSITION AND METHODS OF
TITLE OF INVENTION: TREATING USING THE SAME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
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APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 STATE: MISSINGER STATE: MISSINGER STATE: MISSINGER STATE: MISSINGER STATE: United States

COUNTRY: United States

COMPUTER: ENDABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
CURRENT APPLICATION DAAPA:
APPLICATION NUMBER: US/07/791,213D
FILING DATE: 13-NOV-1991
CLASSIFICATION NUMBER: UP 2-306745
FILING DATE: 13-NOV-1990
ATTORNEY/AGENT INFORMATION:
NAME: Meuth, Donna M
REGISTRATION NUMBER: 36,607
REFERENCE/DOCKET NUMBER: 029650-032
TELEPHONE: (703) 836-6620
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       75.0%; Score 9; DB 1;
100.0%; Pred. No. 0;
Live 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                       E: Burns, Doane, Swecker & Mathis P.O. Box 1404
                                                                                                                                                 Sequence 39, Application US/07791213D Patent No. 5409895 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 9, Application US/08133804
Patent No. 5534254
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEFAN. (703) 836-2021
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Query Match
Best Local Similarity 100.0
Matches 1; Conservative
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                                                                                                                                                                                                                                                                                                                                                                       NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Burns, Dox
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 MOLECULE TYPE: peptide
                                                                                                                                                                                                                                                                                                                                                                                                                                              STREET: P.O. Box
CITY: Alexandria
STATE: Virginia
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TOPOLOGY: linear
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APPLICANT: Huston
                                                                                                                            US-07-791-213D-39
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APPLICANT: 1
APPLICANT: 1
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TITLE OF INVENTION: Imaging
WHERE POSTENCES: 11
CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 ; OTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)" US-08-133-804-9
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ZIP: 5317-1914
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OOMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.30
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/354,240A
FILING DATE: 12-DEC-1994
CLASSIFICATION: 435
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/133,804
FILING DATE:
CLASSIFICATION: 424
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Sequence 12, Application US/08354240A
Patent No. 56/0356;
GENERAL INFORMATION:
APPLICANT: Sherf, Bruce A.
TITLE OF INVENTION:
NUMBER OF SEQUENCES: 20
CORRESPONDENCE ADDRESS:
ADDRESSEE: Dewitt Ross & Stevens, S.C.
STRRET: 8000 Excelsior Drive, Suite 401
CITY: Madison
STATE: WI
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             DB 1;
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                                                                                                                                                                                                                                                                                                                                                                                                         COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ATTORNEY AGENT INFORMATION:
NAME: Relley; Robin D.
REGISTATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054,
TELECOMMUNICATION INFORMATION:
TELEFAN: 617-248-7477
TELEFAX: 617-2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Conservative
                                                                                                                                                                                                                         CITY: Boston
STATE: Massachusetts
COUNTRY: USA:
ZIP: 02109
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               TOPOLOGY: linear MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     NAME/KEY: Protein LOCATION: 1..2
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Query Match
Best Local Similarity
Matches 1; Conserva
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GENERAL INFORMATIES.

APPLICANT: KNAWAORI, TOSHIOTI
APPLICANT: KNAWAORI, TOSHIOTI
APPLICANT: KNAWAORI, TOSHIOTI
APPLICANT: KNAWAORI, TOSHIOTI
APPLICANT: NOBUHARA, MASAHIOD
TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE SAME
TITLE OF INVENTION: INHIBITION PROCESS, DRG COMPOSITION AND METHODS OF
TITLE OF INVENTION: TREATING USING THE SAME
TITLE OF INVENTION: TREATING USING THE SAME
CORRESPONDENCE ADDRESS:
ADDRESSEE: BJRNS, DOANE, SWECKER & MATHIS
STREET: P.O. BOX 1404
CITY: Alexandia
STREET: P.O. BOX 1404
CITY: Alexandia
CONTRY: United States
STREET: P.O. BOX 1404
COMPUTER: ELOPPM;
MEDIUM TYPE: FloppW disk
COMPUTER: ELOPPM;
MEDIUM TYPE: PLOPPM;
MEDIUM NUMBER: JP - 2306745
FILING DATE: JR - NOV-1990
PRIOR APPLICATION NUMBER: JP - 2306745
FILING DATE: JR - NOV-1990
PRIOR APPLICATION NUMBER: JP - 2306745
FILING DATE: JR - NOV-1990
PRIOR APPLICATION NUMBER: JP - 6607
REFERENCE/DOCKET NUMBER: JF - 6607
REFERENCE/DOCKET NUMBER: JF - 6607
PELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
TELECOMMUNICATION INPORMATION:
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                               Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           75.0%; Score 9; DB 1
ilarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
                                                    Pred. No. 0;
Mismatches
                            75.0%; Score 9;
100.0%; Pred. No
ttive 0; Mismat
                                                                                                                                                                                                                                                                                   Sequence 23, Application US/08293150A Patent No. 5792629 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    RESULT 7
US-08-293-150A-39
: Sequence 39, Application US/08293150A
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEFAX: (703) 836-2021
INFORMATION FOR S2Q ID NO: 23:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acid
TYPE: amino acid
           Query Match
Best Local Similarity 100.v
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                ; MOLECULE TYPE: peptide US-08-293-150A-23
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Query Match
Best Local Similarity
'-has 1; Conservat
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APPLICANT: Huston, James S.
APPLICANT: Houston, L. L.
APPLICANT: Houston, L. L.
APPLICANT: Houston, L. L.
APPLICANT: Houston, L. L.
APPLICANT: Hig, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For
TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 11
CORRESPENDENCE ADDRESS:
ADDRESSED: Testa, Hurwitz & Thibeault/Patent Department
STREET: Exchange Place, 53 State Street
CITY: Boston
                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ; LOCATION: 1..2
; OTHER INFORMATION: /note- "C-Terminal Tail (Ser-Cys)"
US-08-461-838-9
                                                                                                                                                                                                                                                                                                                                                                                  Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/461,838
                                                                                                                                                                                                                                                                                                                                                                              Query Match 75.0%; Score 9; DB 1; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches
ATTORNEY/AGENT INFORMATION:
NAME: Sara, Charles S.
REGISTATION NUMBER: 34506.029
REFERENCE/DOCKET 1000
TELECOMMUNICATION INFORMATION:
TELEPHONE: 608-831-2100
INFORMATION FOR SEQ ID NO: 12:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               RESULT 5
US-08-461-838-9; Sequence 9, Application US/08461838; Patent No. 5753204
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CLASSIFICATION: 424
ATTORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054,
TELECOMMUNICATION INFORMATION:
TELEPHOR: 617-248-7477
TELEFAX: 617-248-740
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 2 amino acids
                                                                                                                                                                                                                                                      ; TOPOLOGY: linear
; MOLECULE TYPE: protein
US-08-354-240A-12
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    TYPE: amino acid
STRANDEDNESS: single
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      TOPOLOGY: linear
MOLECULE TYPE: protein
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Massachusetts
                                                                                                                                                                                                                                        TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ZIP: 02109
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Massachusetts
: USA
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MOLECULE TYPE: protein
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STATE: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    COUNTRY: U.S.A.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               US-08-465-380-307
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                                                                                                               TITLE OF INVENTION: POLYPEPTIDE, DNA FRAGMENT ENCODING THE TITLE OF INVENTION: SAME AND PROCESS FOR PRODUCING THE SAME, AND ENZYME TITLE OF INVENTION: INHIBITION PROCESS, DRUG COMPOSITION AND METHODS OF TITLE OF INVENTION: TREATING USING THE SAME NUMBER OF SEQUENCES: 110
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Gaps
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APPLICANT: Oppermann, Hermann
APPLICANT: Oppermann, Hermann
APPLICANT: Houston, L. L.
APPLICANT: Ring, David B.
TITLE OF INVENTION: Biosynthetic Binding Proteins For
TITLE OF INVENTION: Imaging
NUMBER OF SEQUENCES: 1
CORRESPONDENCE ADDRESS:
ADDRESSEE: Testa, Hurwitz & Thibeault/Patent Department
STREET: Exchange Place, 53 State Street
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                CUITE Alexander
STATE: Virginia
STATE: Virginia
COUNTRY: United States
ZIP: 22313-1404
COMPUTER READABLE FORM:
MEDIUM TYPE: Floppy disk
COMPUTER: IBM PC compatible
COMPUTER: Patentin Release #1.0, Version #1.25
SOFTWARE: Patentin Release #1.0, Version #1.25
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/293,150A
FILING DATE: 19-AUG-1994
FILING DATE: 19-AUG-1994
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                DB 1;
                                                                                                                                                                                                                               CORRESPONDENCE ADDRESS:
ADDRESSEE: BURNS, DOANE, SWECKER & MATHIS
STREET: P.O. Box 1404
CITY: Alexandria
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FILING DATE: 19-AUG-1994

PRICASIFICATION: 514

PRICASIFICATION: 514

PRICASIFICATION DATA: 05/791,213

PILING DATE: 13-NOV-1990

PRICASIFICATION NUMBER: US 07/791,213

PRICASIFICATION NUMBER: US 2306745

FILING DATE: 13-NOV-1990

ATTORNEY/AGENT INFORMATION:

NAME: Meuth, Donna M.

REGISTRATION NUMBER: 36,607

REFERENCE/DOCKET NUMBER: 029650-049

TELECOMMUNICATION INFORMATION:

TELECOMMUNICATION 186-6620

TELECHAX: (703) 836-6620

TELECHAX: (703) 836-6201

INFORMATION FOR SEQ ID NO: 39:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                75.0%; Score 9;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Sequence 9, Application US/08461386 Patent No. 5837846
                                             APPLICANT: MORISHITA, Hideaki
APPLICANT: KANAMORI, TOShinori
APPLICANT: NOBUHARA, MASAHIRO
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ; MOLECULE TYPE: peptide US-08-293-150A-39
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   linear
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Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    GENERAL INFORMATION:
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Patent No. 5792629
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 FOPOLOGY:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                US-08-461-386-9
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APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Tris H.L. Mensens, Marc J. Lauwereys,
APPLICANT: Tris G.J. Gansemans, Matthew Moyle,
APPLICANT: Peter W. Bergum
TITLE OF INVENTION: REMATODE-EXTRACTED ANTICOAGULANT
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   ; NAME/KEY: Protein
; LOCATION: 1...2
; OTHER INFORMATION: /note= "C-Terminal Tail (Ser-Cys)"
US-08-461-386-9
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Length 2;
COMPUTER READABLE FORM:
MEDIUM TYPE: RIDAPPY disk
COMPUTER: IBM PC compatible
OPERATING SYSTEM: PC-DOS/MS-DOS
SOFTWARE: PatentIn Release #1.0, Version #1.25
APPLICATION NUMBER: US/08/461,386
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          DB 2;
0;
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100.0%; Pred. No. 0;
tive 0; Mismatches
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COMPUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 APPLICATION NUMBER: US/08/465,380 FILING DATE: June 5, 1995 CLASSIFICATION: 530
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Sequence 307, Application US/08465380 Patent No. 5863894
                                                                                                                                                     FILING DATE:
CLASSIFICATION: 424
ATORNEY/AGENT INFORMATION:
NAME: Kelley, Robin D.
REGISTRATION NUMBER: 34,637
REFERENCE/DOCKET NUMBER: 2054
TELECOMMUNICATION INFORMATION:
TELEPHONE: 617-248-7477
TELEFAX: 617-248-7477
TELEFAX: 617-248-7100
INFORMATION FOR SEQ ID NO: 9:
SEQUENCE CHARACTERISTICS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                              LENGIH: 2 amino acids
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                                                                                                                                                                                                                                                                                                                                                                                                                                                  TYPE: amino acid
STRANDEDNESS: single
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Gaps

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Gaps
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APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Joris H.L. Marco J. Lauwereys,
APPLICANT: Yannick G.J. Gansemans, Matthew Moyle,
APPLICANT: Yannick G.J. Gansemans, Matthew Moyle,
APPLICANT: Peter W. Bergum
APPLICANT: ABARTON:
APPLICANT: ADRESS:
ADDRESSEE: Lyon & Lyon
STREET: Suite 4700
CITY: Los Angeles
APPLICANT: California
                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        OTHER INFORMATION: Xaa in location 2 is an OTHER INFORMATION: amino acid.
                                                                                                                                                                                                                                                     FEATURE:
OTHER INFORMATION: Xaa in location 2 is an
CHER INFORMATION: amino acid.
US-08-486-397-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                            DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                         75.0%; Score 9; DB 2
100.0%; Pred. No. 0;
iive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                COMPUTER: IEM Compatible
COMPUTER: IEM COMPATIBLE
SOFTWARE: IEM COMPATIBLE
SOFTWARE: WORD PREfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,399
FILING DATE: June 5, 1995
CLASSIPICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: OCLOBER 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REGISTRATION NUMBER: 30,158
REGISTRATION NUMBER: 31,270
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELEPHONE: (213) 489-1600
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               COUNTRY: U.S.A.
ZIP: 90071
ZOMUTER READABLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: Storage
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Sequence 307, Application US/08486399 Patent No. 5866543
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FRAGMENT TYPE: internal fragment FEATURE:
                                                                                                                                                                                  TOPOLOGY: linear FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
TYPE: amino acids
              TELEX: 67-3510
INFORMATION FOR SEQ ID NO:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Conservative
                                                                                                                                                          TYPE: amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                            Query Match
Best Local Similarity
Matches 1; Conservi
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          linear
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   APPLICANT: George P. Vlasuk, Patric H. Stanssens, APPLICANT: George P. Vlasuk, Patric H. Stanssens, APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys, APPLICANT: Yves R. Laroche, Laurent S. Jespers, APPLICANT: Yannick G.J. Gansemans, Matthew Moyle, APPLICANT: Peter W. Bergum
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 357
CORRESPONDENCE ADDRESS:
ADDRESSEE: Lyon & Lyon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  DB 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Xaa in location 2 is an amino acid.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               75.0%; Score 9; DB 2
llarity 100.0%; Pred. No. 0;
Conservative 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
MEDIUM TYPE: 184 Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: US/08/486,397
FILING APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING APPLICATION NUMBER: 18, 1994
ATTORNEY AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
RECISTRATION NUMBER: 30,158
REFERENCE/DOKET NUMBER: 213/269
TELECOMMUNICATION INFORMATION:
MEDIUM TELECOMMUNICATION INFORMATION:
MEDIUM TELECOMMUNICATION INFORMATION:
MEDIUM TYPE: ALL AND ALL A
PRIOR APPLICATION DATA:

APPLICATION NUMBER: 08/326,110
FILING DATE: 0ctober 18, 1994
ATTONNEY AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 213/268
TELEPHONE: (213) 489-1600
TELEPHONE: (213) 955-0440
TELEXA: (213) 955-0440
TELEXA: (213) 955-0440
TELEXA: (213) 955-040
TELEXA: (213) 955-0440
TELEX: 2 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Sequence 307, Application US/08486397 Patent No. 5866542 GENERAL INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ADDRESSEE: Lyon & Lyon STREET: 633 West Fifth Street STREET: Suite 4700
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (213) 955-0440
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             STREET: Suite 4700 CITY: Los Angeles STATE: California COUNTRY: U.S.A. ZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              OTHER INFORMATION:
COTHER INFORMATION:
US-08-465-380-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Query Match
Best Local Similarity
Matches 1; Conserv
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RESULT 13
US-08-634-641-307
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                                                                                                                   Gaps
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                                                                                                                                                                                                                                                                                                                   Sequence 307, Application US/08461965
Patent No. 5872098
GENERAL INFORMATION:
GENERAL INFORMATION:
APPLICANT: George P. Vlasuk, Patric H. Stanssens,
APPLICANT: Joris H.L. Mensens, Marc J. Lauwereys,
APPLICANT: Yaves R. Larcche, Laurent S. Jespers,
APPLICANT: Yanick G.J. Gansemans, Matthew Moyle,
APPLICANT: Peter W. Bergum
TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT
TITLE OF INVENTION: PROTEIN
NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
                                                                                                                0; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 0; Indels
                                                                     Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CTHER INFORMATION: Xaa in location 2 is an CTHER INFORMATION: amino acid. US-08-461-965-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 75.0%; Score 9; DB 2; Best Local Similarity 100.0%; Pred. No. 0; Matches 1; Conservative 0; Mismatches
                                                                   75.0%; Score 9; DB 2;
100.0%; Pred. No. 0;
tive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            COUNTRY: US.A.

ZIPALE:
ZIPI: 90071
ZIPI: 90071
ZOMBUTER READABLE FORM:
MEDIUM TYPE: 3.5 Diskette, 1.44 Mb
MEDIUM TYPE: 3.5 Diskette, 1.44 Mb
MEDIUM TYPE: Storage
COMPUTER: IBM Compatible
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Mord Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION NUMBER: 08/08/461,965
FILING DATE: June 5, 1995
CLASSIFICATION: 530
PRIOR APPLICATION DATA:
APPLICATION NUMBER: 08/326,110
FILING DATE: October 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 210/243
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION INFORMATION:
TELECOMMUNICATION SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       ADDRESSEE: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
                                                                                                                Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             CITY: Los Angeles
STATE: California
                                                                   Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               linear
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US-08-486-399-307
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Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Length 2;
                                                                                                                                                                            APPLICANT: Laroche, Yves Rene APPLICANT: Laroche, Yves Rene APPLICANT: Jespers, Laurent Stephane APPLICANT: Gansemans, Yannick Georges Jozef APPLICANT: Moyle, Matthew APPLICANT: Bergum, Peter W. TITLE OF INVENTION: NEMATODE-EXTRACTED ANTICOAGULANT TITLE OF INVENTION: PROTEIN CORRESPONDENCE ADDRESS:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               75.0%; Score 9; DB 2;
100.0%; Pred. No. 0;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               ) OTHER INFORMATION: Xaa in location 2 is ; OTHER INFORMATION: amino acid. US-08-634-641-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb MEDIUM TYPE: storage
COMPUTER: Storage
COMPUTER: Storage
COMPUTER: BW COMPATIBLE
OPERATING SYSTEM: IBM P.C. DOS 5.0
SOFTWARE: Word Perfect 5.1
CURRENT APPLICATION DATA:
APPLICATION DATA:
APPLICATION NUMBER: US/08/634,641
FILING DATE: APRLICATION DATA:
APPLICATION NUMBER: PCT/US95/1323
FILING DATE: October 17, 1995
APPLICATION NUMBER: 08/486,397
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/465,380
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
FILING DATE: June 5, 1995
APPLICATION NUMBER: 08/461,965
APPLICATION NUMBER: 08/461,965
APPLICATION NUMBER: 08/461,965
APPLICATION NUMBER: 08/326,110
FILING DATE: October 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REGISTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 219/136
                                                                          APPLICANT: Vlasuk, George P. Vlasuk
PAPLICANT: Stanssens, Patrick Eric Hugo
APPLICANT: Mensens, Joris Hilda Lieven
APPLICANT: Lauwereys, Marc Josef
Sequence 307, Application US/08634641
Patent No. 5955294
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FRAGMENT TYPE: internal fragment FEATURE:
                                                                                                                                                                                                                                                                                                                                                                            STREET: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Annual
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
TYPE: amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          ELECOMMUNICATION INFORMATION
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            CITY: Los Angeles
STATE: California
COUNTRY: U.S.A.
ZIP: 90071
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Query Match
Best Local Similarity
Matches 1; Conserv
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                                                          GENERAL INFORMATION:
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Indels
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INHIBITORS AND ANTICOAGULANT PROTEIN
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   75.0%; Score 9; DB 3;
100.0%; Pred. No. 0;
ive 0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    CTHER INFORMATION: Xaa in location 2 is OTHER INFORMATION: amino acid. US-09-249-471-307
                                   NUMBER OF SEQUENCES: 356
CORRESPONDENCE ADDRESS:
ADDRESSES: Lyon & Lyon
STREET: 633 West Fifth Street
STREET: Suite 4700
CITY: Los Angeles
STREET: California
COUNTRY: 0.5.A.
ZIP: 90071
COMPUTER READBLE FORM:
MEDIUM TYPE: 3.5" Diskette, 1.44 Mb
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       5, 2001, 12:01:54
                                                                                                                                                                                                                                                                                                                                                                                                          FILING DATE:
PRIOR APPLICATION DATA:
PRIOR APPLICATION NUMBER:
APPLICATION NUMBER:
FILING DATE:
COCTOBER:
APPLICATION NUMBER:
PCT/US95/13231
FILING DATE:
OCTOBER:
APPLICATION NUMBER:
O8/486,399
FILING DATE:
June 5, 1995
APPLICATION NUMBER:
PLING DATE:
APPLICATION NUMBER:
PLING DATE:
June 5, 1995
APPLICATION NUMBER:
O8/465,380
FILING DATE:
June 5, 1995
APPLICATION NUMBER:
O8/461,965
FILING DATE:
June 5, 1995
APPLICATION NUMBER:
O8/461,965
FILING DATE:
June 5, 1995
APPLICATION NUMBER:
O8/326,110
                                                                                                                                                                                                                                                                                                     COMPUTER: IBM Compatible OPERATING SYSTEM: IBM P.C. DOS 5.0 SOFTWARE: Word Perfect 5.1
                                                                                                                                                                                                                                                                                                                                                                                            US/09/249,471
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 216/270
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      FILING DATE: OCTODER 18, 1994
ATTORNEY/AGENT INFORMATION:
NAME: BIGGS, SUZANNE L.
REJESTRATION NUMBER: 30,158
REFERENCE/DOCKET NUMBER: 216/27
TELECOMMUNICATION INFORMATION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   FRAGMENT TYPE: internal fragment
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       TELEPHONE: (213) 489-1600
TELEFAX: (213) 955-0440
TELEX: 67-3510
INFORMATION FOR SEQ ID NO: 307:
SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Ouery Match
Best Local Similarity 100.
                                                                                                                                                                                                                                                                                                                                                               CURRENT APPLICATION DATA:
APPLICATION NUMBER: US
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Search completed: February
Job time: 336 sec
TITLE OF INVENTION:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TOPOLOGY:
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                                                                                                                                     Sequence 52, Application US/08818253

Patent No. 5998204

GENERAL INFORMATION:
APPLICANT: Trien, Roger Y.
APPLICANT: Miyawaki, Atsushi
TITLE OF INVENTION: FLUORESCENT PROTEIN SENSORS FOR
TITLE OF INVENTION: DETECTION OF ANALYTES
NUMBER OF SEQUENCES: 61
CORRESPONDENCE ADDRESS:
ADDRESSEE: Fish & Richardson P.C.
STREET: 4225 Executive Square, Suite 1400
CITY: La Jolla
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             APPLICANT: Moyle, Matthew
APPLICANT: Bergum, Peter W.
TITLE OF INVENTION: NEMATODE-EXTRACTED SERINE PROTEASE
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Length 2;
                                                                                                                                                                                                                                                                                                                                                                                                          STATE: CA
COUNTRY: USA
ZIP: 92037
COMPUTER READBLE FORM:
MEDIUM TYPE: Diskette
COMPUTER: IBM COMPATIBLE
OPERATING SYSTEM: Windows 95
SOFRWARE: FRACESO for Windows Version 2.0b
CURRENT APPLICATION DATA: WS/08/818,253
FILING DATE: 14-MAR-1997
PRIOR APPLICATION NUMBER:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Vlasuk, George Phillip
Stanssens, Patrick Eric Hugo
Messens, Joris Hilda Lieven
Lauvereys, Marc Josef
Laroche, Yves Rene
Jespers, Laurent Stephane
Gansemans, Yannick Georges Jozef
Moyle, Matthew
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            FILING DATE:
ATTORNEY, AGENT INFORMATION:
NAME: Haile, Ph.D., Lisa A.
REGISTRATION NUMBER: 38,347
REFERENCE/DOCKET NUMBER: 07257/043001
TELECOMMUNICATION INFORMATION:
TELEPHONE: 619/678-5070
ITELEFAX: 619/678-509
INFORMATION FOR SEQ ID NO: 52:
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Pred. No. 0;
0; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    75.0%; Score 9; 100.0%; Pred. N
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Sequence 307, Application US/09249471
Patent No. 6040441
GENERAL INFORMATION:
APPLICANT: Vlasuk, George Phillip
APPLICANT: Stanssens, Patrick Erit
APPLICANT: Messens, Joris Hilda Li
APPLICANT: Lauwereys, Marc Josef
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             SEQUENCE CHARACTERISTICS:
LENGTH: 2 amino acids
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              MOLECULE TYPE: peptide US-08-818-253-52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        amino acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Query Match
Best Local Similarity
Matches 1; Conserv
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  RESULT 15
US-09-249-471-307
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              TOPOLOGY:
                                                                                                                            US-08-818-253-52
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APPLICANT:
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Gaps